AR241: 4, AR282: 4, AR192: 4, AR241: 4, AR282: 4, AR241: 4, AR247: 4, AR312: 4, AR312: 4, AR312: 4, AR273: 4, AR243: 4, AR273: 4, AR243: 4, AR299: 3, AR198: 3, AR299: 3, AR300: 3, AR299: 3, AR299: 3, AR299: 3, AR299: 3, AR299: 2, AR297: 1, AR299: 1, AR289:	S0052: 2 H0550: 1, H0428: 1 and	50052: 1.	S0052. 1 S0052. 1	S0052: 1	S0052: 1	S0052: 1	S0052: 1	S0052: 2, S0222: 1, L0455: 1, S0036: 1 and L0366: 1.
•						Met-1 to Met-14.	Met-23 to Val-30.	
	4270	4271	4272	4273	4274	4275	4276	4277
	14 - 100	155 - 250	23 - 115	101 - 199	229 - 339	129 - 233	116 - 208	147 - 335
	1673	1674	1675	1676	1677	1678	1679	1680
	826717	564574	824145	8663998	826451	663997	604485	825345
	HNGIU16	HNGIX91		-	-		+	HNGJL07
·	1663	1664	1665	1666	1667	1998	1669	16/0

L0794: 7, L0768: 2, S0052: 2, H0040: 1, L0789: 1 and S0216: 1.	S0052: 1	AR060: 6, AR316: 5, AR089: 3 H0673: 1 and S0428: 1.	AR060: 9, AR316: 7,		S0428: 1	AR089: 32, AR316: 24,	AR060: 17	71	AR060: 3, AR316: 3,	AK089: 1 S0428: 1	AR060: 7 AR316: 6	. 4	L0745: 9, L0548: 1, S0002:	1, S0428: 1, S0432: 1 and	L0746: 1.	S0053: 1 and S0028: 1.	1 0760: 3 1 0/30: 3 10/676.	2, H0333: 2, L0794: 2, L0783:	2, S0053: 2, S0328: 2, L0748:	2, L0759: 2, H0265: 1, H0556:	1, S0430: 1, H0657: 1, H0341:	1, S0360: 1, S0132: 1, H0393:	1, L0717: 1, H0486: 1, S0010:	1, H0318: 1, H0052: 1, L0471:	1, S0051: 1, H0038: 1, H0040:	1, H0412: 1, H0494: 1, S0016:	1, H0130: 1, S0210: 1, S0422:	 1, L0774: 1, L0655: 1, L0527:	1, L0657: 1, L0789: 1, S0216:	1, L0438: 1, L0352: 1, S0330:	1, S0146: 1, S0032: 1, L0749:
Gly-25 to Asn-37.			Phe-33 to Leu-38,	Pro-45 to Gln-58,	Ala-81 to Phe-86, Gly-88 to Gln-97.	Pro-29 to Tyr-36.											Ser-78 to Asn-34							,							
4278	4279	4280	4281			4282			4283		4284					4285	4286	2021													
49 - 162	251 - 418	313 - 396	86 - 481			290 - 445			403 - 453		299 - 463					69 - 173	77 - 176	2										-			
1681	1682	1683	1684			1685		, 6, ,	1686		1687					1688	1689														
608638	825336	836060	899408			835021		0,0,00	830002		834936					609892	654859	)													
	HNGJU60	HNGKB09	HNGKW35			HNGKY94		000 10101	HINGLD28		HINGOY36					$\dashv$	HNHAC43				•										
1671	1672	1673	1674			1675		1	0/01		1677				1	1678	1679														

						103950, 120580,	131440, 139130,	142680, 176260,	190450, 200990,	216950, 600228,	600414, 600618,	960209																				
						12p13																										
1, L0779: 1, L0777: 1, L0757: 1, S0242: 1, S0276: 1 and H0423: 1.	AR089: 24, AR316: 18, AR060: 12 S0053: 1	S0053: 2, H0171: 1 and S0037: 1	H0313: 1 and S0053: 1.	H0313: 1 and S0053: 1.	S0053: 1		AKU6U: 8	H0556: 34, H0265: 17,	S0422: 15, L0/42: /, L0///:	7, H0013: 6, H0156: 6, L0776:	6, H0547: 6, L0439: 6, L0471:	5, L0805: 5, H0144: 5, H0624:	4, H0141: 4, H0585: 4, H0657:	4, H0599: 4, H0560: 4, L0517:	4, H0519: 4, L0779: 4, S0134:	3, S0376: 3, H0619: 3, H0351:	3, S0278: 3, S0222: 3, H0486:	3, H0457: 3, L0163: 3, L0455:	3, H0040: 3, L0770: 3, L0772:	3, L0375: 3, L0747: 3, L0731:	3, L0599: 3, H0423: 3, H0170:	2, H0583: 2, S0007: 2, L0717:	2, H0586: 2, H0497: 2, S0010:	2, H0263: 2, H0050: 2, H0024:	2, H0356: 2, H0615: 2, H0634:	2, H0616: 2, S0426: 2, L0369:	2, L0809: 2, S0378: 2, L0748:	2, L0759: 2, H0595: 2, H0542:	2, H0171: 1, S0114: 1, S0218:	1, H0656: 1, S0116: 1, H0662:	1, H0305: 1, H0589: 1, H0459:	1, H0638: 1, S0358: 1, S0132: 1, H0369: 1, H0431: 1, H0250:
		Ala-24 to Tyr-29.																														
	4287	4288	4289	4290	4291	4292						-																		·		
	54 - 137	157 - 282	344 - 349		162 - 275	705 - 722																		•								
	1690	1691	1692	1693	1694	1695																										
	612845	604993	006609	604996	668254	535713																										
	HNHAD34	HNHAG83	HNHAH06	HNHAJ65	HNHAL61	HNHAP58																										
	1680	1681	1682	1683	1684	1685													·									•. •				

1, L0021: 1, H0002: 1, H0575: 1, H0196: 1, H0581: 1, S0049: 1, H0196: 1, H0178: 1, H0123: 1, H0041: 1, H0178: 1, H0123: 1, H0057: 1, H0373: 1, H0428: 1, H0039: 1, H0644: 1, T0067: 1, H0272: 1, H0623: 1, T0041: 1, H0272: 1, H0529: 1, L0648: 1, L0535: 1, L0769: 1, L0646: 1, L0535: 1, L0769: 1, L0646: 1, L0769: 1, L0509: 1, L0792: 1, L0569: 1, L0509: 1, L0792: 1, L0665: 1, S0053: 1, H0698: 1, H0520: 1, H0436: 1, S0328: 1, S0330: 1, H0436: 1, S0328: 1, S0330: 1, H0436: 1, R0390: 1, L0755: 1, L0758: 1, H0645: 1, L0758: 1, L0758: 1, H0445: 1, S0434: 1, S0436: 1, L0795: 1 and H0422: 1.	S0053: 1 S0053: 1 S0053: 1 AR182: 4, AR269: 4, AR060: 3, AR184: 3, AR296: 3, AR282: 3, AR298: 3, AR285: 3, AR298: 3, AR285: 3, AR298: 3, AR241: 3, AR299: 2, AR241: 3, AR299: 2, AR291: 2, AR299: 2, AR291: 2, AR299: 2, AR291: 2, AR299: 2, AR296: 2, AR268: 2, AR266: 2,
	4293 4294 4295
	214 - 252 144 - 236 777 - 887
	1696 1697 1698
	603386 825316 704789
	HNHAW34 HNHAW35 HNHAY26
	1686

		104770, 107300, 107670, 110760, 126133	145001, 146790,	159440, 159440,	159440, 173610, 186780, 191030,	191315, 600923,	601412, 601518,	001632, 002491													
		1q22-q24																			
AR258: 2, AR096: 2, AR267: 2, AR232: 2, AR218: 2, AR263: 2, AR277: 2, AR237: 2, AR175: 2, AR231: 2, AR238: 2, AR265: 2, AR229: 1, AR233: 1, AR299: 1, AR277: 1, AR033: 1	H0052: 1, S0053: 1 and S0216: 1.	7, AR204: 5, AR282:	AR206: 4, AR053: 4, AR186: 4, AR273: 4,		AR060: 4, AR294: 3, AR313: 3, AR183: 3,	3, AR265:	3, AR061:	AK182: 3, AK298: 3, AR296: 3 AR205: 3	3, AR243:	3, AR277:	AR309: 3, AR269: 3,	 	• •	• • •	AK283: 2, AK213: 2, AB270: 2 AB202: 2		íα	•	2, AR089:	4	AR039: 2, AR267: 2,
	4296	4297														-	•				
	99 - 113	243 - 356																			
	1699	1700																			
	866180	839696	,																		
	HNHAY86	HNHAZ20													-						
	1689	1690																			

AR234: 2, AR232: 2, AR096: 2, AR231: 1, AR284: 1, AR175: 1, AR310: 1, AR229: 1, AR227: 1 S0053: 1	S0052: 1 and S0053: 1.	AR089: 48, AR316: 35, AR060: 24	S0132: 9, S0046: 6, S0418: 5,	H0617: 5, H0050: 4, L0602: 4,	L0/51: 4, H0295: 3, H0013: 3,	LU/63: 3, LU/68: 3, LU639: 3, 1 0747: 3, 1 0506: 3, 50040: 2	H0294: 2, S0420: 2, S0376: 2,	S0360: 2, S0045: 2, H0250: 2,	H0266: 2, H0622: 2, H0551: 2,	H0623: 2, H0494: 2, L0646: 2,	L0363: 2, L0767: 2, L0649: 2,	L0775: 2, L0518: 2, L0783: 2,	S3014: 2, S0028: 2, L0588: 2,	L0581: 2, L0595: 2, S0276: 2,	H0556: 1, H0685: 1, H0341: 1,	 <u>,                                    </u>	<u>_</u>	<u>_</u>	<b>ન</b> .	<b>_</b> ;	<u>.</u>	<u>.</u>		H0606: 1, H0212: 1, H0135: 1,	H0090: 1, H0040: 1, H0634: 1,	H0264: 1, H0268: 1, H0412: 1,	H0056: 1, S0386: 1, L0564: 1,	S0440: 1, H0509: 1, S0210: 1,	S0002: 1, L0637: 1, L0772: 1,	L0372: 1, L0645: 1, L0764: 1,
	Val-12 to Lys-18.	His-4 to Thr-10, Ser-16 to Ser-25	Leu-107 to Ser-121,	Pro-143 to Asn-152.																			•							
	4298	4299														•														
	395 - 529	214 - 672																												
	1701	1702																												
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L0533: 1, L0378: 1, L0655: 1, L0807: 1, L0383: 1, L0809: 1, L0791: 1, L0666: 1, L0663: 1, L0664: 1, L0665: 1, S0053: 1, H0144: 1, H0547: 1, H0593: 1, H0435: 1, H0658: 1, H0670: 1, H0672: 1, S0152: 1, S0332: 1, H0479: 1, L0749: 1, L0748: 1, L0754: 1, L0749: 1, L0779: 1, L0755: 1, L0758: 1, L0601: 1, S0011: 1, H0668: 1, H0665: 1, H0422: 1, S0424: 1 and H0506: 1	S0053: 1 and S0216: 1	1	H0341: 1, S0001: 1 and S0053: 1.	S0428: 1 and S0053: 1	S0053: 1 and H0422: 1.	S0053: 1	AR060: 6, AR316: 4,	AR089: 3 S0046: 1 and S0053: 1.	AR089: 20, AR316: 17,	AR060: 14 S0053: 2 and S0046: 1.	S0053: 1	L0439: 7, S0222: 6, H0013:	6, L0771: 4, L0805: 3, L0776:   3, S0053: 3, 1, 0438: 3, 1, 0755:	3, S0007: 2, H0156: 2, S6028:	2, H0555: 2, L0366: 2, H0394:	1, H0556: 1, S6024: 1, H0657:	1, L0005: 1, H0443: 1, S6016:	1, H0486: 1, H0244: 1, H0250:	1, H0253: 1, S0010: 1, H0363: 1 H0264: 1 H0272: 1 1 0351:	1, L0370: 1, L0638: 1, L0794:
			Ser-18 to Met-41, Gly-58 to Glu-69.		Asp-6 to Gln-12.		1	7	Asn-25 to Thr-33.	<del>-</del>			9 6	<u>, w</u>			-		<b>-</b>	1
	4300	4301	4302	4303	4304	4305	4306		4307		4308	4309				-				
	657 - 665	161 - 301	374 - 601	293 - 361	55 - 207		226 - 264		73 - 192		115 - 231	32 - 127								
	1703	1704	1705	1706	1707	1708	1709		1710		1711	1712								
	604924	566816	566866	694657	651348	606791	664496		634691		737905	824072								
	HNHBG18	$\dashv$	_		$\dashv$	-	HINHCT22		HNHCT47		-+	HNHCZ54								
	1693	1694	1695	1696	1697	1698	1699		1700		1701	1702								

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1, L0775: 1, L0659: 1, S0328: 1, S0152: 1, H0576: 1, S0028: 1 and H0543: 1	AR060: 8, AR316: 7, AR089: 5 S0222: 1 and S0053: 1	S0053: 1	AR060: 5, AR316: 4,	AK089: 2 H0494: 17, H0551: 13,	S0418: 12, H0521: 11, S0358:	6, L0659: 6, L0526: 6, L0665: 6, C0442: 5, H0510: 5, S0226	0, 30442. 3, IIO319. 3, 30328. 5, H0545. 4, H0553. 4, H0509.	4, L0794: 4, L0766: 4, L0666:	L0663: 4, L0603: 4, H0543:	4, H0265: 3, S0356: 3, H0617:	3, H0488: 3, L0761: 3, L0667:	L0662: 3, L0809: 3, S0126:	S0406: 3, L0759: 3, H0294:	T0049: 2, H0657: 2, S0354:	S0376: 2, S0444: 2, S0360:	S0408: 2, H0587: 2, H0486:	H0375: 2, H0687: 2, H0181:	H0412: 2, T0042: 2, H0561:	H0529: 2, L0638: 2, L0771:	2, L0657: 2, L0664: 2, S0053:	L0438: 2, H0520: 2, H0547:	H0522: 2, H0555: 2, L0740:	L0/31: 2, L0/33: 2, L0/31: 1 0757: 3, 50434: 3, 1,0505:	1 0581: 2, 30434: 2, 10330:	2, 20001: 2, 20079: 2, 110024: 1 1 H0556: 1 H0685: 1 H0713.	1, 110250: 1, 110065: 1, 110/13:	024: 1,110050: 1, f10464:	1, HU001: 1, HU003: 1, SU348:	1, S0420: 1, S0410: 1, H0637:	1, H0722: 1, L0717: 1, H0437:	1, H03/0: 1, H0586: 1, H0559:	l, H0250: 1, H0427: 1, H0156: 1, H0599: 1, T0048: 1, H0581:
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	207 - 299	140 - 184	272 - 586																							_						
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, H0251: 1, H0546: , N0006: 1, L0471: , H0620: 1, H0024: , L0163: 1, H0083: , H0060: 1, H0266: , H0688: 1, T0006: , H0124: 1, H0708: , H0688: 1, L0640: , L0575: 1, L0800: , L0773: 1, L0768: , L0557: 1, L0661: , L0557: 1, L0661: , L0551: 1, L0689: , H0693: 1, H0689: , H0683: 1, H0689: , H0683: 1, H0689: , H0659: 1, L058: , L059: 1, L058: , L0752: 1, L058: , L0752: 1, L0593: , S0932: 1, S0027: , L0752: 1, L0593: , S0026: 1, H0653: , H0423: 1, S0424:	13,					13, 12, 10,	9,
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			Glu-33 to Tyr-38.	Leu-24 to Gly-32.			
	4313	$\Box$	十	4316 L	4318	4319	
	1175 - 1201	94 - 186	19 - 198	291 - 395	311 - 358	332 - 433	
•	1716	1717	1718	1720	1721	1722	
	738043	610113	610114	664497	610094	607867	
	HNHDI17	HNHDL37	HINHDM21 HNHDP57	+-	$\vdash$	HNHDW34	
	1706	1707	1700	1710	1711	1712	

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AP178. 0 AP770. 8	8, AR180:		 6, AR262:	6, AR053:	AR312: 6, AR245: 6,	AR183: 6, AR299: 6,	6, AR269:	6, AR274:	6, AR176:	AR204: 5, AR182: 5,	5, AR181:	5, AR268:	5, AR297:	5, AR175:	5, AR225:	4, AR212:		4, AR196:		4, AR174:	4, AR234:	4, AR23/:	3, AK213:	3, AK233:	3, AR239:	3, AR291:	3, AR282:	AR295: 3, AR189: 3,
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AR290: 3, AR255: 3, AR203: 3, AR104: 3, AR227: 3, AR200: 3, AR170: 3, AR177: 3, AR217: 2, AR311: 2, AR224: 2, AR190: 2, AR250: 2, AR232: 2, AR055: 2, AR263: 2, AR061: 2, AR188: 2, AR061: 2, AR283: 2, AR172: 2, AR205: 1, AR171: 1, AR210: 1	S0053: 1	S0053: 1	S0053: 1	S0053: 1	S0053: 1	S0053: 1 and L0745: 1.	S0053: 1	S0053: 1	S0053: 1	AR176: 6, AR162: 5, AR161: 5, AR163: 5.					AR244: 3, AR273: 3,					AR052: 3, AR200: 3,
	Arg-29 to Ser-35, Pro-39 to Gly-45.						Pro-34 to Arg-43.	Met-1 to Pro-10.		Gln-93 to Cys-99.	7	7	7	7	4	7	7	1	,	7
	4320	4321	4322	4323	4324	4325	4326	4327	4328	4329										
	170 - 322	282 - 350	282 - 287	180 - 230	70 - 132	7 - 111	219 - 371	267 - 371	88 - 207	67 - 612										
	1723	1724	1725	1726	1727	1728	1729	1730	1731	1732										
	608145	569774	608299	825295	668256	610115	638122	638123	607421	854712										
	HNHDX28	Н		_	HNHEF49	HINHEF70	HNHEG30	HNHEH38	HNHEL22	HNHEN70										
	1713	1714	1715	1716	1717	1718	1719	1720	1721	1722										

					AR309:	3. AR182:	3,		
2, AR282: 2, AR275: 2, AR275: 2, AR173: 2, AR180: 2, AR297: 2, AR286: 2, AR266: 2, AR265: 2, AR266: 2, AR268: 2, AR288: 2, AR278: 1, AR288: 2, AR164: 2, AR288: 2, AR288: 2, AR288: 2, AR288: 1, AR289: 2, AR189: 2, AR172: 2, AR189: 1, AR184: 1, AR283: 1, AR283: 1, AR289:					AR223:		2,		
2, AR257; 2, AR204: 2, AR173: 2, AR180: 2, AR297: 2, AR266: 2, AR265: 2, AR265: 2, AR268: 2, AR268: 2, AR268: 2, AR288: 2, AR288: 2, AR288: 1, AR288: 1, AR189: 2, AR189: 2, AR189: 1, AR184: 1, AR184: 1, AR281: 1, AR281: 1, AR281: 1, AR281: 1, AR289:			<del>.</del> .		AR170:		2,		
2, AR275. 2, AR130. 2, AR180. 2, AR266. 2, AR266. 2, AR265. 2, AR265. 2, AR268. 2, AR261. 2, AR288. 2, AR288. 2, AR288. 2, AR288. 2, AR288. 2, AR164. 2, AR172. 2, AR172. 2, AR189. 2, AR189. 1, AR283. 1, AR283. 1, AR289.	-				AR201:		2,	_	
2, AR204: 2, AR173: 2, AR287: 2, AR266: 2, AR266: 2, AR265: 2, AR267: 2, AR261: 2, AR261: 2, AR273: 2, AR172: 2, AR172: 2, AR172: 2, AR172: 1, AR283: 1, AR283: 1, AR283: 1, AR289: 1, AR289: 1, AR289: 1, AR289: 1, AR290: 1, AR291:					AR205:				
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		Ser-20 to Ser-26.			Met-1 to Pro-6, Leu-56 to Gln-68.					Arg-19 to Arg-29.		<del>,</del>		Asn-25 to Thr-33.	1	$\mathcal{H}$	Gln-17 to Asn-28.	Met-1 to Ser-6.	<del>.</del>	<u> </u>	<u> </u>	9
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	1748	1749	1750	1751	1752	1753	1754	1755					1756
	658743	855956	604903	533057	777856	800877	839255	853372					838255
	HNHHA47	HNHHN22	HNHHW53	HNHIB40	HNHKI74	HINHKV56	HNHLD80	HNHLS76				:	HNHLZ47
	1738	1739	1740	1741	1742	1743	1744	1745					1746

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	Pro-22 to Gly-29, His-48 to Ser-62.		Glu-25 to Ser-37.			Met-1 to Gin-12.	Gln-14 to Ser-32, Pro-53 to Leu-60.	Pro-20 to Pro-28, Pro-36 to Cys-41.
	4354	4355	4356	4357	4358	4359	4360	4361
	51 - 242	220 - 330	25 - 192	250 - 351	255 - 392	17 - 112	539 - 739	384 - 527
	1757	1758	1759	1760	1761	1762	1763	1764
	836065	834974	838256	836066	836067	840374	838264	802009
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	1747	1748	1749	1750	1751	1752	1753	1754

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	82 - 231 4362	178 - 1254 4363
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HODDG72	HODDJ25	HODDN21	НОББОЗ1	900000
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	4487	4488	4489	4490
	251 - 346	166 - 384	173 - 349	25 - 174
	1890	1891	1892	1893
	598716	566793	580867	599378
	НРОДО17	нРООО28	HPDDT14	HPEAA65
	1880	1881	1882	1883

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4491	4492	4493	4494
207 - 314	533 - 643	189 - 293	53 - 109
1894	1895	1896	1897
637097	753957	638144	605010
HPEAG24	HPEBA84	HPEBF91	HPEBI09
1884	1885	1886	1887

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H0331: 1 and H0166: 1.	AR060: 8, AR316: 7, AR089: 7 L0805: 5, L0750: 3, L0777: 3, L0803: 2, L0804: 2, H0520: 2, H0547: 2, L0744: 2, L0779: 2, L0752: 2, L0731: 2, S0282: 1, R0360: 1, H0411: 1, H0042: 1, H0581: 1, H0251: 1, H0596: 1, H0123: 1, H0520: 1, H0014: 1, H0169: 1, S0422: 1, L0776: 1, L0791: 1, H0519: 1, H0521: 1, H0522: 1, L0780: 1, L0753: 1 and L0756: 1, L0759: 1, L0759: 1, L0756: 1, L0759: 1, L075	S0050: 1, H02/1: 1, H0169: 1 and S0052: 1.	AR089: 34, AR316: 27, AR060: 22 L0758: 7, L0748: 4, L0731: 4, L0604: 4, H0728: 3, H0486: 3, H0457: 3, H0169: 3, H0543: 3, S0360: 2, L0659: 3, H0543: 3, S0360: 2, L0655: 2, L0809: 2, L0666: 2, H0725: 2, S0328: 2, L0759: 2, S0434: 2, H0542: 1, L0785: 1, H0713: 1, H0716: 1, L0785: 1, H0484: 1, S0420: 1, H0688: 1, H0428: 1, H0708: 1, S0366: 1, H0409: 1, H0413: 1, S0366: 1, L0766: 1, L0657: 1, L0653: 1, L0766: 1, L0657: 1, L0663: 1, H0684: 1, S0013: 1, L0757: 1, H0665: 1, S0196: 1 and S0398: 1.
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	4495	4496	4497
	1 - 33	78 - 173	973 - 1002
	1898	1899	0061
	774564	581083	829318
	HPFCJ75	HPFCP75	HPFDB66
	1888	1889	1890

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824856	638164	747704	852658	847429
HPFDD28	HPFDI47	HPIAF35	HPIAK27	HPIAL55
1891	1892	1893	1894	1895

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3 3 3 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	Thr-4 to His-10.	Asn-54 to Thr-61.  5 2 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Ser-6 to Gly-11.
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	801924	161988	840379
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	588944							843567	835506	841653														
	HPIBI40							HPIBT19	HPJAA82	HPJAB75														
	1900							1901	1902	1903													•	

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	854893	826719	812766	608307
	HPJAN76	HPJAU94	HPJAW78	HPJBS16
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1919	1920	1921	1923
798101	743232	886192	852869
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	559369						638147	658715	639092			
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	1930 772 - 804	1931 166 - 315
	753435	651354
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	HPMDD27
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	Pro-18 to Gly-23.			
	4530	4531	4532	4533
	58 - 177	233 - 328	172 - 267	749 - 865
	1933	1934	1935	1936
	638148	567303	795709	756931
	HPMDF45	HPMDP57	HPMEG72	HPMFM70
,	1923	1924	1925	1926

			600281, 600281
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·		Thr-31 to Gly-36, Ala-39 to Asp-45, Tyr-65 to Gly-71, Arg-79 to Gly-88.	
4534	4535	4536	4537
166 - 288	95 - 151	67 - 333	12 - 50
1937	1938	1939	1940
597457	844865	590991	830452
HPMFP48	HPMFW01	HPMGM06	HPMGW43
1927	1928	1929	1930

	174810, 600624, 600993, 602080	
	18q21.1- 18q21.2	
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		His-48 to Ser-61, Ala-66 to Val-72.
	4538	4539
	110 - 277	1351 - 1593
	1941	1942
	822864	900362
	HPMGY89	HPMKB09
	1931	1932

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			Ser-28 to Phe-39.				Gln-20 to Leu-25.	Gln-20 to Leu-25.	
	4541	4542	4543	4544	4545	4546	4547 4548	4549	4550
·	57 - 152	85 - 204	233 - 355	164 - 298	284 - 292		21 - 116 251 - 361	251 - 361	234 - 248
	1944	1945	1946	1947	1948	1949	1950	1952	1953
	610023	853550	782957	788813	740261	716371	612818	867512	559372
	HPMSH96	HPQAJ25	HPQAJ27	HPQAN50	<b>НР</b> QАО80	HPQAW27	НРОВС90 НРОВЈ48	нРQВ148	HPQBL67
	1934	1935	1936	1937	1938	1939	1940 1941	1942	1943

AR060: 1 S0136: 136	AR060: 6, AR316: 4, AR089: 2 S0136: 85	S0136: 2	AR060: 6, AR316: 5, AR089: 3 S0136: 73 and L0684: 1.	S0136: 115, L0731: 10, H0624: 9, L0803: 9, S0026: 9, S0358: 7, L0439: 7, S0418: 6,	L0803: 0, L0734: 0, H0019: 3, L0659: 5, L0666: 5, L0750: 5, L0756: 5, L0770: 5, H0170: 4	S0360: 4, H0553: 4, L0770: 4,	L0662: 4, H0547: 4, L0759: 4, H0351: 3, H0486: 3, H0013: 3,	H0046: 3, H0050: 3, L0471: 3,	H0200: 3, H0023: 3, L0774: 3, L0776: 3, H0144: 3, H0520: 3,	S0380: 3, L0777: 3, S0040: 2,	H0295: 2, H0661: 2, S0420: 2, S0376: 2, H0722: 2, S0476: 2	H0369: 2, H0592: 2, H0333: 2,	H0574: 2, H0427: 2, H0575: 2,	H0051: 2, S0022: 2, H0615: 2, H0090: 2, S0438: 2, S0440: 2,	L0598: 2, L0769: 2, L0766: 2,	L0518: 2, L0663: 2, H0726: 2,	S0126: 2, H0539: 2, H0521: 2,	H0696: 2, H0555: 2, L0740: 2,	L0747: 2, L0753: 2, L0755: 2,	L0757: 2, L0593: 2, H0686: 1,	S0212: 1, H0484: 1, H0671: 1,	H0662: 1, S0356: 1, S0442: 1,	S0354: 1, S0444: 1, H0645: 1, S0222: 1 H0441: 1 H0409: 1
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1954	1955	1956	1957	1958																•			
826413	684368	566799	655361	802969																			-
HPQBT17	нРQСF94	нРQС162	HPQRS74	HPRAD30																			
1944	1945	1946	1947	1948																			

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H0586: 1, S0280: 1, H0599: 1, H0599: 1, H0375: 1, H0375: 1, H0375: 1, H0378: 1, H0328: 1, H0328: 1, H064: 1, L0809: 1, H064: 1, S0454: 1, H0647: 1, H0647: 1, H0667: 1, H0667: 1,	AR228: AR211: AR231: AR299: AR299: AR299: AR299: AR297: AR275: AR217: AR275:
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	9. <del>4.</del> -, -,	l: 1,	3: 1	9: 11,	9, L0	0:5, F	0: 4, I	): 4, L	4: 3, S	5: 3, L	3.8	, 2 S	5.2.1		2 .	2,1	; ; ∃	, c	, c , c	; ; ; ; ;	; —; ; —; ; —	2: 1. H	: 1, S(	: 1, SC		<u>_</u>	: 1, H	I: 1, H	: 1, H	:: 1, H	: 1, H			: 1, H(	: 1, H(	:. 1, U	: 1, H	: 1, H
	1, AR039: 1. AR104:	l, AR311:	1, AR163:	L0757: 13, L0759: 1	L0485: 10, L0731: 9, L0747:	6, H0013: 5, T0010: 5, H0031	H005	4, L0764: 4, L0740: 4, L0750:	L0758: 4, H0014: 3, S0366:	L0662: 3, L0666: 3, L0438:	L0756; 3, L0599; 3, S0376;	2, S0046: 2, H0619: 2, S0010	2, H0032; 2, H0135; 2, 1,0598;	1.0761: 2, 1.0646: 2, 1.0766:	L0803: 2, L0775: 2, L0806:	L0776: 2, 1.0657: 2, 1.0659	1,066	H0547: 2, E0003; 2, IIO144;	H0521.2 1 0439.2 1 0751.	10749.2 10777.2 10592.	L0362: 2. H0170: 1. H0265	1, H0556: 1, H0222: 1, H0662	S0418: 1, L0005: 1, S0356:	S0358: 1, S0360:	H0489: 1, H0729:	S0468: 1, H0208:	S0222: 1, H0370: 1, H0587:	H0497: 1, H0331: 1, H0486:	T0109: 1, H0156:	H0575: 1, T0082: 1, H0706:	S0346: 1, H0318: 1, H0581	1, H0052: 1, H0309: 1	1, H0178: 1, H0123: 1	1, S0050: 1, L0163: 1	1, S6028: 1, H0687: 1, H0286:	1, H0328: 1, H0622: 1, L0455:	1, H0708: 1, H0316: 1, H0038	, H0551: 1, H0164: 1, H0413.
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<u>ਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑਜ਼ੑ</u>	• AAAA, K.
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			Arg-76 to Lys-91.
	4558	4559	4560
	151 - 306	171 - 263	281 - 607
	1961	1962	1963
	638153	566873	813512
	HPRCF50	HPRCL58	HPRCM72
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	1954

1, S0212: 1, H0663: 1, H0402: 1, S0418: 1, S0420: 1, S0442: 1, S0354: 1, S0376: 1, S0046: 1, H0351: 1, H0411: 1, H0600: 1, H0609: 1, H0156: 1, H0599: 1, H0609: 1, S0346: 1, H0318: 1, H0581: 1, L0040: 1, L0471: 1, L0163: 1, S0388: 1, S6028: 1, H0267: 1, H0687: 1, H0673: 1, L0194: 1, T0006: 1, H0673: 1, L0194: 1, T0006: 1, H0673: 1, L0456: 1, H0316: 1, H0623: 1, L0456: 1, L0598: 1, H0529: 1, L0763: 1, L0598: 1, L0768: 1, L0646: 1, L0571: 1, L0768: 1, L0649: 1, L0775: 1, L0806: 1, L0653: 1, L0657: 1, L0515: 1, L0683: 1, R0519: 1, H0593: 1, H0650: 1, H0650: 1, H0660: 1, S0330: 1, S0374: 1, L0565: 1, H0658: 1, L0780: 1, H0436: 1, H0631: 1, L0750: 1, H0436: 1, L0589: 1, L0599: 1, H0136: 1, S0276: 1 and H0422: 1.	H0428: 2, L0369: 2, L0771: 2, L0803: 2, L0774: 2, L0805: 2, L0665: 2, L0731: 2, L0759: 2, H0556: 1, H0402: 1, S0360: 1, L0717: 1, H0013: 1, S0280: 1, L0770: 1, L0769: 1, L0646: 1, L0736: 1, L0790: 1, S0052: 1, L0809: 1, L0790: 1, S0052: 1, L0438: 1, H0520: 1, H0593: 1, H0672: 1, H0539: 1, L0788: 1, S0196: 1 and H0422: 1.
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	25 - 105 4563
	852240 1966
	1956 HPRTH56

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AR089: 27, AR316: 23, AR060: 19 L0769: 5, L0759: 4, H0402: 3, S0376: 3, H0424: 3, L0749: 3, S0360: 2, S0222: 2, L0535:	2, L0774; 2, L0806; 2, L0384; 2, L0666; 2, H0265; 1, S0218; 1, H0484; 1, H0255; 1, S0354; 1, S0007; 1, H0392; 1, H0559; 1, H0486; 1, T0039; 1, T0060; 1, L0021; 1, H0004; 1, H0524; 1, H0052; 1, H0204; 1, H0024; 1, H00	1, H0405: 1, H0617: 1, H0606: 1, H0090: 1, H0272: 1, H0494: 1, S0002: 1, L0770: 1, L0772: 1, L0372: 1, L0771: 1, L0768: 1, L0794: 1, L0381: 1, L0775: 1, L0789: 1, L07	1, H0547: 1, H0659: 1, H0658: 1, H0672: 1, H0521: 1, H05716: 1, L0744: 1, L0779: 1, L0758: 1, H0445: 1, H0423: 1 and H0422: 1	AR060: 22, AR316: 16, AR089: 10 H0618: 3, H0619: 2, H0424: 2, L0776: 2, H0136: 2, S0276: 2, H0716: 1, H0261: 1, H0250: 1, H0575: 1, H0036: 1, H0610: 1, H0401: 1, H0169: 1, H0616: 1, L0351: 1, H0652: 1, L0768: 1, L0766: 1, L0803: 1, L0659: 1, S0126: 1, H0672: 1, H06539: 1, S0406: 1, H0672: 1, H0539: 1, L0779: 1, L0777: 1 and	AR226: 36, AR232: 21, AR061: 14, AR238: 13,
Gly-35 to Ser-40, Ser-61 to Arg-72, Pro-148 to Arg-172.				Pro-19 to Pro-26, Gly-33 to Ser-38, Gly-45 to Tyr-52, Ser-65 to Ser-71.	Met-1 to Pro-7.
4564				4565	4566
102 - 665				266 - 481	149 - 262
1967				1968	1969
884167				655362	838154
HPTRE80				HPTR142	HPTRL95
1957				1958	1959

AR196: 12, AR060: AR228: 11, AR089: AR239: 11, AR237: AR161: 10, AR163: AR231: 10, AR163: AR233: 8, AR234: AR231: 9, AR166: AR233: 8, AR234: AR201: 7, AR182: AR203: 6, AR229: AR204: 6, AR209: AR205: 5, AR181: AR300: 5, AR288: AR206: 5, AR288: AR207: 5, AR188: AR207: 5, AR188: AR208: 5, AR188: AR208: 5, AR209: AR209: 4, AR209: A

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	4567	
	224 - 325	
	1970	
	655363	
	HPTRQ52	
	1960	

176100, 176100, 178300, 230000, 255800	
25.00.5.3.4.5.5.2.5.2.5.5.3.3.3.4.5.5.5.5.5.5.5.5.5.5.5.5.5.5	6. 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
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7, L0754: 6, H0265: 5, 5, L0758: 5, S0420: 4, 4, L0517: 4, H0657: 3, 3, H0052: 3, H0083: 3, H0082: 3, H0082: 3, H0052: 3, H0049: 3, H0124: 3, H0295: 2, L0747: 2, H0641: 2, H0641: 2, L0764: 2, L0774: 2, L0775: 2, L0776: 2, L0776: 2, L0776: 2, L0776: 2, L0755: 2, L0595: 2, L0595: 2, L0755: 2, L0595: 2, L0756: 1, H0661: 1, H0662: 1, H0662: 1, H0662: 1, H0661: 1, H0673: 1, H067	H0616: 1, H0087: H0264: 1, T0041: H0647: 1, S0002: L0642: 1, L0652: 1, L0664: 1, H0144: H0593: 1, H0690: H0659: 1, H0658: H0651: 1, H0539: S0406: 1, H0576:
L0731: 7, L0754: 6, H0265: 5, H0556: 5, H0556: 5, L0758: 5, S04206: 4, S0408: 3, H0556: 3, L0758: 3, H0657: 3, H0658: 3, H0653: 3, H0653: 3, H0653: 3, L0768: 3, L0768: 3, L0768: 3, L0768: 3, L0768: 3, L0768: 2, H0771: 2, H0678: 2, H0778: 2, H0678: 2, L0778: 2, L0768: 2, H0678: 1, H0658: 1, H0668: 1, H0668: 1, H0668: 1, H0668: 1, H0668: 1, H0668: 1, H0678: 1, H0678	H0040: 1, H0616: 1, H0087: T0067: 1, H0264: 1, T0041: 1 H0131: 1, H0647: 1, S0002: 1 L0772: 1, L0642: 1, L0662: 1 L0767: 1, L0657: 1, L0659: 1 L0382: 1, L0664: 1, H0144: 1 S0374: 1, H0593: 1, H0690: 1 H0682: 1, H0659: 1, H0539: 1 H0666: 1, H0651: 1, H0539: 1 H0661: 1, S0406: 1, H0576: 1
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L0743: 1, L0740: 1, L0750: 1, L0779: 1, H0445: 1 and S0436: 1.	L0809: 6, L0751: 5, L0747: 5, H0556: 4, L0749: 4, L0774: 3, L0748: 3, H0265: 2, S0003: 2, L0439: 2, L0757: 2, L0439: 2, L0757: 2, L0439: 2, L0757: 2, L0758: 2, H0664: 1, S0366: 1, H0664: 1, S0376: 1, H0618: 1, H0623: 1, H0618: 1, H0253: 1, H0039: 1, H0609: 1, H0560: 1, H0560: 1, L0769: 1, L5566: 1, L0769: 1, L0760: 1, L0760: 1, L0760: 1, L0760: 1, L0362: 1 and S0276: 1	H0424: 2, L0615: 1, H0213: 1 and H0658: 1.	L0439: 3, L0749: 3, H0551: 2, L076: 2, L076: 2, L076: 2, H0547: 2, L075: 2, H0459: 1, S0418: 1, H0637: 1, L0717: 1, S0222: 1, T0039: 1, L0021: 1, H0253: 1, H0264: 1, H0264: 1, H0264: 1, L0770: 1, L0770: 1, L0748: 1, L0748: 1, L0748: 1, L0748: 1, L0748: 1, L0748: 1, L0759: 1, L0755: 1, L0731: 1, L0759: 1, L0755: 1, L0731: 1, L0759: 1, H0352: 1.	S0356: 3, H0052: 3, H0457: 3, S0142: 3, H0265: 2, S0358: 2, H0253: 2, H0581: 2, H0424: 2, L0769: 2, L0761: 2, L0764: 2, L0666: 2, H0144: 2, H0672:
		Arg-22 to Trp-27, Gln-29 to Glu-36.	Pro-27 to Cys-33.	
	4568	4569		4571
	80 - 142	210 - 326		/66 - 8/3
	1971	1972	1973	19/4
	566875	612865	561954	831983
	HPTTH35	HPTTI65	HPTTT62	HP1 V H24
	1961	1962	1963	1964

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		141750, 141800, 141800, 141800, 141800, 141850, 141850, 141850, 141850, 141850, 156850, 186580, 191092, 600140, 600273, 601313, 601785	
	1	16p13.3	16
2, H0521: 2, L0750: 2, L0596: 2, L0591: 2, L0591: 2, H0556: 1, S0040: 1, H0657: 1, S0212: 1, H0638: 1, S0418: 1, S0240: 1, S0418: 1, S0240: 1, H0507: 1, H0607: 1, H0502: 1, L0021: 1, H0575: 1, T0048: 1, H0421: 1, H0641: 1, H0641: 1, H0641: 1, L0637: 1, L0800: 1, L0773: 1, L0364: 1, L0804: 1, L0773: 1, L0634: 1, L0659: 1, L0773: 1, L0734: 1, L0659: 1, H0666: 1, S0374: 1, H0658: 1, H0658: 1, H0650: 1, S0378: 1, S0378: 1, S0378: 1, S0378: 1, L0740: 1, L0743: 1, L07	H0424: 15, S6024: 1, H0033: 1, H0213: 1, L0769: 1 and L0805: 1.	L0764: 4, L0439: 4, H0424: 3, L0659: 3, H0617: 2, L0666: 2, L0756: 2, L0757: 2, H0543: 4, H00343: 1, H0034: 1, H0036: 1, H0036: 1, L0668: 1, L0438: 1, H0682: 1, H0436: 1, L0777: 1, L0778: 1 and H0423: 1.	L0809: 5, L0779: 5, L0758: 5, L0769: 4, L0764: 4, L0777: 4, H0634: 3, L0748: 3, L0750: 3, L0731: 3, S0222: 2, L0163: 2, L0763: 2, L0772: 2, L0775: 2, L0776: 2, H0651: 2, L0749:
	Pro-24 to Gln-44.	Ser-23 to Arg-30, Glin-42 to Pro-48, Arg-50 to Arg-56.  1 1 1 1 1 1 1 1 1	5 3 3 2 2
	4572	4573	4574
	144 - 278		271 - 291
	1975	1976	1977
	695753	897503	636064
	HPTVH59	HPTVI04	HPTVI96
	1965	1966	1967

	115660, 139250, 148500, 150200, 154275, 162100, 170500, 170500, 170500, 170500, 230200, 249000, 253250		602134
	17q24		2p24.3-p24.1
2, L0759: 2, H0657: 1, S0376: 1, S0300: 1, L0622: 1, H0427: 1, H0009: 1, L0471: 1, T0006: 1, H0424: 1, H0674: 1, H0708: 1, L0771: 1, L0768: 1, L0794: 1, L0766: 1, L0774: 1, L0375: 1, L0806: 1, L0789: 1, H0648: 1, H0521: 1, S0406: 1 and H0506: 1.	AR060: 6, AR316: 5, AR089: 4 L0439: 4, L0803: 3, H0590: 2, L0483: 2, H0163: 2, L0805: 2, S0374: 2, H0658: 2, H0696: 2, H0717: 1, S0116: 1, S0358: 1, L0717: 1, H0052: 1, H0194: 1, H0184: 1, L0738: 1, H0545: 1, S0316: 1, S0003: 1, S0364: 1, S0316: 1, S0036: 1, H0572: 1, L0638: 1, L0766: 1, L0776: 1, L0789: 1, L0664: 1, L0665: 1, L0789: 1, L0664: 1, L0776: 1, H0710: 1, H0521: 1, S0013: 1, S0406: 1, L0744: 1, L0752: 1, L0758: 1, L0605: 1, S0026: 1 and H0543: 1.	L0776: 2, L0748: 2, H0170: 1, H0685: 1, H0580: 1, H0052: 1, S6028: 1, L0771: 1, L0775: 1, H0520: 1, S0013: 1, L0749: 1, L0777: 1 and H0543: 1.	L0439: 17, L0770: 8, L0438: 8, L0740: 6, L0749: 6, L0803: 5, L0748: 5, L0755: 5, L0766: 4, L0731: 4, S0374: 3, H0670: 3, H0685: 2, S0444: 2, H0431: 2, H0494: 2, H0538: 2, L0659: 2, L0791: 2, L0777: 2, L0747: 2, L0779: 2, L0777: 2, L0747: 2, L0777: 2, L0747: 2, L0777: 2, L0777
		1	8 4 5 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
	4575	4576	4577
	74 - 148		311 - 352
	1978	1979	7980
	783074	601692	50195
	HPVAA15	HPVAB20	HPVAB03
	1968	1969	0/61

2, H0170: 1, H0556: 1, S0342: 1, H0713: 1, S0402: 1, L04144: 1, S0110: 1, S0001: 1, H0662: 1, S0356: 1, S0358: 1, S0356: 1, S0356: 1, S0356: 1, S0356: 1, H0629: 1, H0619: 1, H0647: 1, H0619: 1, L0769: 1, L0	1 1	H0036: 1 L0749: 3, L0748: 2, H0581:	S0346: 1, S0049: 1, H0124: 1, H0144: 1, S0044: 1, L0748: 1 and I 0747: 1	S0051: 1, S0044: 1 and H0423: 1.
		Thr-16 to Ala-27.		
	4578	45/9	4581	4582
	91 - 210	144 - 263 46 - 171	305 - 409	83 - 262
	1981	1982	1984	1985
	750951	732157	536008	747520
	HPVAF86	HPWA089	HPWAS27	HPWAT86
	1971	1973	1974	1975

L0775: 2, L0638: 1, L0803: 1, S0044: 1, L0751: 1, L0756: 1, L0759: 1 and H0668: 1	AR089: 14, AR316: 12, AR060: 9 S0044: 2 and L0803: 1	AR089: 52, AR316: 31, AR069: 18, L0747: 18, H0656: 13, H0436: 10, H0271: 9, L0751: 9, S0474: 8, H0581: 8, H0179: 8, H0650: 7, L0662: 7, L0756: 7, H0650: 7, L0662: 7, L0756: 7, H0550: 5, H0050: 5, H0050: 5, H0050: 5, H0050: 5, H0050: 5, H0050: 4, H0590: 4, H0590: 5, H0171: 4, H0390: 4, H0591: 4, L0771: 4, H0390: 4, H0591: 4, L0671: 3, H0662: 3, H0651: 4, H0739: 4, H0749: 3, L0769: 3, H0679: 3, H0679: 3, L0769: 3, H0679: 3, L0769: 3, L0769: 3, H0679: 3, L0769: 3, L0769: 3, L0769: 3, L0779: 3,
		Val-80 to Leu-92, Ser-98 to Lys-104, Pro-111 to Pro-122.
4583	4584	4585
401 - 439	83 - 157	283 - 675
1986	1987	1988
830084	840380	843700
HPWAV82	HPWBA36	HPWTF23
1976	1977	1978

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T H	Val-80 to Leu-92, A Ser-98 to Lys-104, A			S	<u>jilij</u>	<u>н</u>	<u> </u>	<u> </u>	1	<u> </u>	S	S	<u> </u>	1	<u> </u>	<u>, 11</u>	H	<u>1</u>	I	<u> </u>	1	<u> </u>	1	1	S	S	Щ.	<u> </u>	1	4	8	<u> </u>	<u> </u>	Н.	<i>σ</i> .	7
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		484 - 777 4587
,		844737 1990
		1980 HPWTF53

## ISSINS ... ISSIST

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	Lys-22 to Gly-29.	Ala-22 to Asp-33.							Asn-19 to Ser-25.					Thr-36 to Trp-44.	Pro-2 to Gln-7.									,			Pro-26 to Gly-34.						
	4588	4589						000,	4390	4591				4592	4593											4594	4595						
		42 - 152						0	0 - 03	274 - 453				103 - 252	54 - 149				-							95 - 205	3201 - 3320					-	
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	175 - 189	224 - 331	268 - 420	207 - 353	150 - 215
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	621384	823344	621381	562230	637647
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		Arg-7 to Arg-14, Tyr-38 to Glu-43.																																
	4601	4602		4603	4604		4605			4606				4607			4608																	
	124 - 216	304 - 444		147 - 221	87 - 224		86 - 223			248 - 364				281 - 334			235 - 369					•												
	2004	2005		2006	2007		2008		0000	5003				2010			2011				•		•		•									
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	HRACI39	HKADU15		HRDAH04	HRDBA20		HRDBD32		Tagan.	HKDBL01			203 100	HKDDM85		CCSCION	HKDDS22			- ,,											-			
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	Phe-18 to Ala-23.
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	67 - 126	1193 - 1318	51 - 155	75 - 155	74 - 487
	2013	2014	2015	2016	2017
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			HRDFT83	HRGCA01	HRGCA06
	2003	2004	2005	2006	2007

	1pter-p12			
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-	Gly-24 to Gly-36, Gly-48 to Arg-57, Ala-61 to Gln-66, Lys-82 to Lys-98.		Ser-21 to Phe-32.	
	4615	4616	4617	4618
	50 - 370	158 - 223	·	254 - 331
	2018	2019	2020	2021
	898233	795038	610614	732148
	HRGSE38	HRLAT43	HRLME03	HROAN20
	2008	2009	2010	2011

									176830, 176830.	182601, 229800,	002134										****			
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L0373: 1.	AR060: 8, AR316: 7, AR089: 6 H0316:1	AR060: 838, AR316: 546, AR089: 233 H0316: 1	AR060: 1 H0316: 1	AR089: 11, AR316: 11, AR060: 11	H0581: 1, H0598: 1 and L0662: 1.	AR060: 7, AR316: 6, AR089: 4	H0395: 1, L0762: 1, L0775:	T0008: 1	AR089: 26, AR316: 21,	AR060: 17 1.0758:15-1.0731:13	L0752: 12, L0755: 9, L0766:	8, H0641: 7, S0474: 6, H0038:	6, H0521: 6, L0439: 6, H0179:	5, L0662: 5, L0655: 5, L0748:	3, LU/ //: 5, LU/59: 5, H0638: 4 H0581: 4 H0581: 4	4, L0809; 4, L0742; 4, L0754.	4, L0749: 4, L0757: 4, S0222:	3, L0666: 3, L0663: 3, S0053:	3, H0522: 3, H0436: 3, L0740:	3, L0750: 3, S0442: 2, H0580:	2, H0156: 2, L0021: 2, S0010:	2, H0014: 2, S0003: 2, S0214:	2, H0031: 2, H0090: 2, H0623:	2, S0002: 2, L0598: 2, L0764:   2, L0768: 2, L0774: 2, L0806:
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	181 - 288	152 - 247	242 - 337	108 - 281	395 - 529	320 - 364	82 - 174	266 - 361	959 - 1087							-			_				_	
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and H0251: 1.	AR060: 7, AR316: 6, AR089: 5 S0410: 9, L0794: 4, L0779: 4, S0360: 3, L0752: 3, L0759: 3, H0295: 2, H0545: 2, L0742: 2, L0747: 2, H0713: 1, H0294: 1, S0114: 1, H0484: 1, H0483: 1, H0661: 1, H0402: 1, S0420: 1, S0358: 1, S0444: 1, S0007: 1, H041: 1, H0536: 1, S0499: 1, H0617: 1, H0334: 1, S0440: 1, L0770: 1, L0769: 1, L0803: 1, L0804: 1, L0809: 1, L0791: 1, R0593: 1, H0660: 1, S0037: 1, S0027: 1, S0028: 1, L0740: 1, L0786: 1, S0106: 1 and	AR089: 5, AR316: 5, AR060: 5 L0731: 17, L0766: 15, H0521: 15, H0457: 13, L0740: 9, H0486: 8, S0358: 6, S0474: 6, L0747: 6, H0423: 6, H0581: 5, L0754: 5, H0638: 4, L0779: 4, H0445: 4, S0134: 3, S0045: 3, S0374: 3, L0757: 3, L0485: 2, S0360: 2, H0586: 2, H0581: 2, H0575: 2, H0560: 2, L0792: 2, L0764: 2, L0662: 2, L0369: 2, L0764: 2, L0665: 2, L03665: 2, L0760: 2, L0768: 2, H0518: 2, H0519: 2, S0328: 2, H0518: 2, H0528: 1, H0170: 1, H0171: 1, T0002: 1, S0114: 1, H0650: 1, H0657: 1, H0656: 1, H0341:
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	823161	866194	600370	603367	800279
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	653 - 694	121 - 270 101 - 190	126 - 212 223 - 243
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	690151	738018	866195
	HSAVZ05	HSAWB58 HSAWH36	HSAWM20 HSAWM74
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	Gly-50 to Asp-57.		Met-1 to Gly-6, Pro-32 to Ser-37, Ser-57 to Gly-62	or or or or or		
	4662	4663	4664	4665	4666	4667
	1363 - 1536	59 - 154	104 - 313	86 - 115	164 - 286	90 - 152
	2065	2066	2067	2068	2069	2070
	872498	606842	598726	606826	566801	604946
	HSAWX70	HSAXC22	HSAXI10	HSAXL49	HSAXL82	HSAXN57
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	Ser-28 to Arg-34.					Lys-33 to His-41.																											
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	260 - 364		158 - 268	20 - 127	151 - 339	139 - 264	1900 - 1971																	-									
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		•	Phe-28 to Arg-33.	Arg-21 to Ser-35.
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	136 - 282		246 - 377	00 - 193
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AR286:	H0271:	7, H0494 5 H0622	3, H0022	4, LU/43 4 H0619	3, H0416	3, H0519		3, H0543	2, H0592	2, T0041:	2, L0667:	2, L0794:	2, L0789:	2, S3014:	2, L0755:	2, L0581:	1, S0342:	1, S0134:	1, H0661	1, S0420:	1, S0132:	1, S0222:	1, H0455;	1, H0632:	1, H0599:	1, S0049:	1, H0204:	1, H0457:	1, H0014:	1, H0188:	1, H0428:	1, H0553:	1, S0366:	1, H0551:	1, L0475:	1, H0625:	1, H0646:	1, L0762:
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Gly-19 to Trp-27, Gln-40 to Trp-45.	Thr-52 to Ile-61, Val-64 to Lys-73.		Ala-61 to Leu-81,	Pro-123 to Glu-131,	Ala-141 to Lys-146.																																
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2112	2113	2114								2115					
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	Gly-1 to Ala-13.		7
	4715	4716	4717
	187 - 468	263 - 373	46 - 111
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	Ala-42 to Pro-47, Pro-62 to His-68, Fro-83 to Asn-92, His-94 to Gln-103, Ala-111 to Asn-119, Thr-131 to Ile-137, Glu-246 to Arg-252, Ser-308 to Cys-315, Lys-326 to Phe-334, Lys-343 to Leu-348.	Arg-21 to Cys-26, Pro-41 to Tyr-47, Val-50 to Lys-58.
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4774	4775	4776	4777	8/
286 - 336	838 - 927	190 - 234		21 - 198
717	2178	2179	2180	7101
600330	784994	617125	786645	7.00740
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2167	2168	2169	2170	

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	590743	838072	562783	737753	600373	830046
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	2240	2242	2243	2244	2245	2246

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AR089: 3 H0038: 1 and L0758: 1.	H0038: 2, S0142: 2, S0152: 2, L0756: 2, L0758: 2, S0424: 2, H0624: 1, H0171: 1, L0393: 1, H0346: 1, H0661: 1, S0376: 1, H0013: 1, H0575: 1, H0123: 1, H0056: 1, S0210: 1, L0648: 1, L0789: 1, L0769: 1, L0664: 1, H0519: 1, L0759: 1, L0608: 1 and S0011: 1.	H0038: 1, H0616: 1 and L0779: 1.	L0748: 8, L0777: 5, L0779: 3, H0046: 2, H0038: 2, L0756: 2, H0057: 1, H0634: 1, H0616: 1, L0662: 1, L0803: 1, L0792: 1, L0666: 1, H0684: 1, H0670: 1 and L0749: 1.	AR089: 11, AR316: 10, AR060: 9 L0438: 7, L0740: 4, H0416: 3, L0803: 3, L0777: 3, S0005: 2, H0544: 2, H0271: 2, H0990: 2, L0766: 2, L0809: 2, H0436: 2, L0748: 2, L0366: 2, H0265: 1, H0556: 1, H0583: 1, H0656: 1, H0576: 1, H0583: 1, H0656: 1, H0341: 1, S0360: 1, S0408: 1, S0132: 1, S0476: 1, H0599: 1, S0222: 1, H0632: 1, H0599: 1, S0049: 1, H0123: 1, H0659: 1, H0553: 1, H0266: 1, H0659: 1, H0553: 1, H0566: 1, H0659: 1, H0561: 1, H0599: 1, H0576: 1, H0659: 1, H0576: 1, H0650: 1, H0576: 1, H0579: 1, H0576: 1, H0579: 1, H0576: 1, H0679: 1, H0576: 1, L0779: 1, L0644: 1, L0771: 1, L0773:
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	4891	4892	4893	4894
	36 - 98		192 - 299	343 - 543
	2294	2295	2296	2297
	637689	824739	823145	862067
	HTEGY81	HTEHB11	НТЕНВ49	HTEHS91
	2284	2285	2286	2287

: 1, L0655: : 1, H0696: : 1, L0780: 422: 1.	16: 1.			: 3, H0591: : 3, L0779: : 2, L0755: : 2, H0170: : 1, S0212: 11, S0046: : 1, H0427: : 1, H0046: : 1, R0250: : 1, H0553: : 1, H0053:	: 1, L0771: : 1, L0519: : 1, H0520: : 1, H0521: : 1, L0747: : 1, L0591:	5: 10, 2, L0759: 1: 1, S0360: 11, H0038: 11, L0803:	16: 1.	i: 37,
1, L0662: 1, L0768: 1, L0655: 1, L0665: 1, H0682: 1, H0696: 1, H0478: 1, L0749: 1, L0780: 1, L0759: 1 and H0422: 1.	H0038: 2 and H0616: 1	H0038: 1	H0038: 2	4, L0749; 4, H0618; 3, H0591; 3, H0038; 3, L0776; 3, L0779; 3, H0644; 2, H0090; 2, L0761; 2, H0693; 2, L0748; 2, L0755; 2, L0759; 2, H0445; 2, H0170; 1, H0650; 1, H0657; 1, S0212; 1, S0420; 1, S0376; 1, S0046; 1, H0549; 1, H0431; 1, H0427; 1, H0050; 1, H0024; 1, S0250; 1, H0615; 1, L0141; 1, H0553; 1, L0142; 1, H0632; 1, H0032; 2, H0032; 3, H0032; 4, H00	1, H0616: 1, H0063: 1, L0771: 1, L0662: 1, L0809: 1, L0519: 1, L0666: 1, H0144: 1, H0520: 1, H0658: 1, S0350: 1, H0521: 1, H0631: 1, S3014: 1, L0747: 1, L0758: 1, L0596: 1, L0591: 1, L0595: 1, S0011: 1 and S0026: 1.	AR089: 10, AR316: 10, AR060: 9 H0616: 3, L0520: 2, L0759: 2, H0294: 1, H0650: 1, S0360: 1, L0021: 1, H0318: 1, H0038: 1, L0475: 1, L0794: 1, L0803: 1, L0779: 1 and L0757: 1.	H0038: 1 and H0616: 1	AR089: 46, AR316: 37, AR060: 29 H0038: 1
								Ser-36 to Ser-41.
	4895	4896	4897			4899	4900	4901
	38 - 130	96 - 185	141 - 257			183 - 257	207 - 227	102 - 227
	2298	2299	2300			2302	2303	2304
	637690	862068	737852			806472	603393	570826
	HTEHV60	HTEHW80	HTEID25			HTEIM62	HTEIV33	HTEIV65
	2288	2289	2290			2292	2293	2294

			91.200																					-							
L0758: 2, H0150: 1, H0038: 1 and L0779: 1.	L0805: 4, L0748: 4, H0038: 3, L0776: 2, L0758: 2, H0170:	1, 50402: 1, 50218: 1, 50444: 1, H0052: 1, H0024: 1, H0032: 1, H0050: 1, 10770: 1, 10662:	1, L0766: 1, L0527: 1, L0558:	1, L0789: 1, S0374: 1, H0593:	l, L0751: 1, L0749: 1, L0779:	1, L0777: 1, L0731: 1, L0608: 1, S0192: 1 and S0412: 1.	AR060: 5, AR316: 5,	AR089: 4	H0169: 6, H0038: 3 and L0665: 1.	H0038: 1	H0038: 1	H0038: 1	L0747: 28, S0192: 21,	L0659: 17, L0740: 12, H0038:	10, L0757: 10, L0758: 10,	S0360: 9, H0545: 9, L0662: 9,	L0775: 9, L0665: 9, L0439: 9,	L0750: 9, L0664: 8, S0358: 7,	H0616: 7, L0666: 7, L0601: 7,	L0777: 6, L0731: 6, L0471: 5,	S0003: 5, H0090: 5, H0547: 5,	S0206: 5, L0748: 5, L0756: 5,	L0752: 5, S0011: 5, S0026: 5,	T0049: 4, L0717: 4, H0575: 4,	H0012: 4, H0620: 4, H0083: 4,	L0770: 4, L0764: 4, L0794: 4,	L0766: 4, L0804: 4, L0375: 4,	L0806: 4, L0754: 4, L0779: 4,	L0596: 4, L0588: 4, L0581: 4,	L0603: 4, H0556: 3, H0431: 3,	H0370: 3, S0010: 3, H0596: 3, L0157: 3, S0388: 3, T0023: 3,
Lys-24 to Asn-29.							7	7		Gln-34 to Lys-40.				<u> </u>		<u> </u>		I	I	1	<u> </u>	<u> </u>	<u> </u>		1		I	I	7		the paral
4902	4903						4904			4905	4906	4907	4908																		
96 - 185	257 - 292						94 - 168			124 - 246	298 - 351	136 - 249	389 - 490																		
2305	2306						2307			2308	2309	2310	2311																		
637691	637687						828178			762831	824733	603409	596826																		
HTEJC50	HTEJD20						HTEJD61			HTEJF31	HTEJI29	HTEJL16	HTEJP65				-														
2295	2296						2297			2298	2299	2300	2301																		

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: 3, L0564 : 3, L0667	3, L0774	: 3, L0809	3, S0122	: 3, L0591	: 2, H068¢	: 2, H065C	2, S0132:	: 2, H0427	. 2. H026	2, H004(	2, T001C	2. H027	2, H064	: 2, H0591	: 2, H0059	2, 1,0369	2, 1,0637	2, 1,0767	2. L0776	2, L0565	: 2, H0670	: 2, H0672	2, H0555	: 2, H0627	2, S0032:	2, L0592	2, L0595:	2, H0170	: 1, T0002	1, H0294	1, S0282:	: 1, H0663	: 1, H0638	1, S0444:	H0208: 1, H0619: 1	1, H0369	: 1, H0587	: 1, H0574
H0031: 3, H0264: 3, L0564: 3, S0372: 3, L0769: 3, L0667: 3,	L0521: 3, L0768: 3, L0774: 3,	L0657: 3, L0783: 3, L0809: 3,	L0663: 3, S0126: 3, S0122: 3,	H0682: 3, L0759: 3, L0591: 3,	H0352: 3, L0353: 2, H0686: 2,	S0040: 2, H0295: 2, H0650: 2,	S0045: 2, S0046: 2, S0132: 2,	H0549: 2, S0222: 2, H0427: 2,	H0156: 2, H0318: 2, H0263: 2,	H0597: 2, H0546: 2, H0046: 2	H0050: 2, H0024: 2, T0010: 2	H0375: 2, H0266: 2, H0271: 2	H0030: 2, H0553: 2, H0644: 2	H0674: 2, H0135: 2, H0591: 2	H0412: 2, H0413: 2, H0059: 2	H0100: 2, S0002: 2, L0369: 2	L0520: 2, L0763: 2, L0637: 2,	L0761: 2, L0372: 2, L0767: 2,	L0649: 2, L0803: 2, L0776: 2,	L0382: 2, L0647: 2, L0565: 2,	H0690: 2, H0659: 2, H0670: 2	H0660: 2, H0648: 2, H0672: 2	S0404: 2, S0406: 2, H0555: 2,	H0478: 2, H0626: 2, H0627: 2	L0070: 2, S3014: 2, S0032: 2,	L0741: 2, L0590: 2, L0592: 2,	.0599: 2, L0593: 2, L0595: 2,	H0543: 2, L0600: 2, H0170: 1	H0171: 1, H0265: 1, T0002: 1	H0140: 1, S0470: 1, H0294: 1	L0785: 1, S0212: 1, S0282: 1	H0484: 1, H0671: 1, H0663: 1	H0664: 1, H0662: 1, H0638:	L0005: 1, S0442: 1, S0444: 1	1, H0208	H0393: 1, S6022: 1, H0369: 1,	H0550: 1, H0455: 1, H0587: 1	H0333: 1, H0331: 1, H0574:
H0031: S0372:	L0521:	L0657:	L0663:	H0682:	H0352:	S0040:	S0045:	H0549:	H0156:	H0597:	H0050:	H0375	H0030:	H0674:	H0412:	H0100:	L0520:	10761:	L0649:	L0382:	H0690:	H0660:	S0404:	H0478:	L0070:	L0741:	L0599:	H0543:	H0171:	H0140:	L0785:	H0484:	H0664:	L0005:	H0676: 1,	H0393:	H0550:	H0333:
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1, H0295: 1, S0418: 1, S0045: 1, S0046: 1, H0619: 1, H0069: 1, S0074: 1, H0046: 1, H0150: 1, S0051: 1, S0051: 1, S0003: 1, H0150: 1, H0615: 1, L0483: 1, H0169: 1, H0212: 1, H0634: 1, H0413: 1, H0100: 1, S0382: 1, S0422: 1, L0776: 1, L0807: 1, L0803: 1, L0666: 1, L0775: 1, L0438: 1, H0519: 1, S0152: 1, S0146: 1, H0436: 1, L0756: 1, L0733: 1, S0434: 1, L0756: 1, L0753: 1, S0434: 1, L0599: 1 and H0422: 1.	H0038: 1	AR089: 168, AR316: 132, AR060: 100	L0758: 6, L0750: 3, L0752:	2, H0619: 1, L0666: 1, L0665:	1, H0144: 1, L0438: 1, L0439:	1, L0747: 1, L0756: 1, L0753:	AR251: 23. AR053	AR310: 21, AR309:		AR213: 18, AR263:	AR313: 16 AR203:	, ,	 _	_	_	AR266: 12, AR241:	_	_		AR282: 10, AR253: AR285: 10 AR033:
		Thr-22 to Gln-35.					Pro-115 to Leu-122,	Lys-164 to Lys-171,	Tyr-307 to Ile-312.											
	4910	4911					4912		_									<u>.</u>		
	41 - 136	457 - 564					634 - 1602					-								
e re	2313	2314					2315						_		•					
	004979	094648					877673													
UTEVNO	UTEVBOY	HIENF82					HTEKV69													
2303	2307	4067					2305													

			103581, 146150, 151670, 203200, 203200, 227220, 600374, 601623, 601780, 601800, 601889, 602117
			15q11.2- 15q22.33
AR096: 9, AR289: 9, AR248: 9, AR240: 8, AR277: 8, AR219: 8, AR284: 8, AR186: 8, AR104: 8, AR291: 8, AR177: 8, AR175: 8, AR296: 7, AR299: 7, AR298: 7, AR218: 7, AR298: 6, AR299: 6, AR298: 6, AR294: 6, AR298: 5, AR297: 5, AR256: 5, AR297: 5, AR256: 5, AR297: 5, AR256: 5, AR297: 5, AR259: 4, AR298: 5, AR298: 5, AR298: 5, AR298: 5, AR298: 5, AR298: 5, AR298: 3, AR298: 5, AR298: 3, AR298: 2, AR298: 3, AR398: 2, AR298: 3, AR398: 3, AR298: 3, AR398: 3, AR298: 3, AR398: 3, AR298: 3, AR398: 3, AR298: 3,	H0038: 1	AR089: 9, AR316: 7, AR060: 5 H0305: 2, H0436: 2, H0581:	AR233: 31, AR293: 20, AR275: 19, AR192: 18, AR182: 16, AR274: 16, AR039: 16, AR298: 15, AR229: 14, AR205: 13, AR292: 13, AR286: 12, AR285: 12, AR204: 12, AR282: 12, AR273: 11,
			Gln-8 to Cys-13, Pro-18 to Leu-24, His-62 to Pro-68.
	4913	4914	4915
	146 - 244	389 - 454	2365 - 2793
	2316	2317	2318
	762832	840031	900824
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	2306	2307	2308

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AR259: 11. A	AR295: 10, A	AR237: 9, A		AR186: 7, A	AR183: 7, A		AR296: 5, A	AR244: 5, A	AR053: 5, AJ			AR179: 3, A	AR284: 3, AJ	AR096: 2, A	AR104: 2, A	AR300: 2, A	AR269: 2, A	AR241: 2, A	AR089: 2, Al	AR033: 1, Al	AR266: 1, Al	H0694: 92, L	8, S0126: 7, L	6, L0776: 6, L	6, L0759: 6, H	5, H0623: 5, L	5, H0656: 4, S	4, H0486: 4, H	4, H0529: 4, L	4, L0655: 4, L	4, S0434: 4, H	3, H0574: 3, H	3, H0644: 3, H	3, L0764: 3, L	3, S0374: 3, H	3, S0330: 3, L	3, S0436: 3, H
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, L0717: 2,	2, 30200. 2, 2. H0032: 2	. S0422: 2,	2, L0768: 2,	2, H0658: 2	S0044: 2, H0555: 2, L0744:	L0751: 2, L0749: 2, L0777:	2, L0758: 2, L0592:	L0731: 2, L0738: 2, L0392: L0581: 2, L0593: 2, H0653:	, LUCCO. 1, TINEA2. 0	S0276: 2, H0543: 2, L0411:	H0556: 1, H0220: 1, S0114:	T0049: 1, H0583: 1, H0650	S0116: 1, H0341: 1, S0212	S0356: 1, S0354: 1, S0444:	S0350: 1, S0354: 1,		, H0619:	1, S0222: 1	1. H0586:	1,10106.1	I, H0036: 1, L0105: 1, 10048:	H0581: 1, H0374: 1, H0046:	, H0123: 1, H0050: 1, L0471	H0355: 1, S0250: 1, H0688	H0029: 1, H0039: 1	T0006: 1, H0553: 1,	1, H0124: 1	H0316: 1, H0591: 1, H0264	H0268: 1, H0413: 1, H0056:	T0042: 1, S0016: 1, H0561:	S0438: 1, H0509: 1, S0150:	H0641: 1, H0646: 1	S0344: 1, S0426: 1,	I, L0521: 1	_		1, L0367: 1	I, H0520: 1	1, H0593: 1, H0689: 1, H0670	I, H0672: 1, S0378: 1, H0518:	, S0152: 1, S0350: 1, H0522
2, S0045: 2, L0717: 2, S0414:	2, F0465: 2, S0260: 2, F0150: 2, S0010: 2, H0032: 2, H0494:	2, S0440: 2, S0422: 2, L0667:	2, L0646: 2, L0768: 2, L0652:	2, L0438: 2, H0658: 2, H0539	2, S0044: 2	2, L0751: 2	2. L0731: 2.	2, L0731: 2 2, L0581: 2	7, LVJ. 01. 2	2, S0276: 2	1, H0556:	1, T0049: 1	1. S0116: 1	1 S0356: 1	1,0000001	1, 30300.	1, S0132: 1	1, H0437: 1	1 H0592:	1, 110022	1, H0030:	1, H0581:	1, H0123:	1, H0355:	1, H0029:	1, T0006: 1	1, H0169: 1	1, H0316:	1, H0268:	1, T0042: 1	1, S0438: 1	1, H0641:	1, S0344: 1	1, L0771: 1, L0521:	1, L0389: 1, L0803:	1, L0775: 1, L0661:	1, L0657: 1, L0367:	1, L0664: 1, H0520:	1, H0593:	1, H0672:	1, S0152: 1
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	824604	762835	851698	834832	762836	740771	732366
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	858755	783141
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	2540	2541	2542

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6, L0662: 5, L0766: 5, L0776: 5, L0748: 5, L0788: 5, L0599: 5, L0764: 4, L0775: 4, S0374: 4, S0206: 4, H0255: 3, S0360: 3, S0408: 3, L0517: 3, L0526: 3, L0749: 3, L0752: 3, L0755: 3, L0759: 3, L0755: 3, L0759: 3, L0759: 3, S0134: 2, H0657: 2, S0418: 2, S0444: 2, L0717: 2, H0013: 2, H0309: 2, H0182: 2, H0182: 2, L0369: 2, L0369: 2, H0182: 2, L0369: 2, L0369: 2, H0182: 2, L0369: 2, H0182: 2, L0369: 2, L03	2, L0770: 2, L0769: 2, L0372: 2, L0641: 2, L0773: 2, L0363: 2, L0806: 2, L0659: 2, H0520: 2, H0547: 2, H0684: 2, H0670: 2, S0380: 2, H0684: 2, L0731: 2, L0743: 2, L0744: 2, L0731: 2, L0757: 2, L0596: 2, L0485: 2, H0506: 2, H0265: 1, H0484: 1, H0661: 1, S0212: 1, H0484: 1, H0661: 1, H0638: 1, R0420: 1, R0486: 1, H0661: 1, H0632: 1, H0486: 1, H0661: 1, H06477: 1, H0699: 1, H0030: 1, H06477: 1, H0599: 1, H0030: 1, H0030: 1, H00477: 1, H0599: 1, H0030: 1,	H0318: 1, H0263: 1, H0596; H0597: 1, H0231: 1, H0545; H0046: 1, L0471: 1, H0024; T0079: 1, H0687: 1, H0428; H0644: 1, H0617: 1, H01553; H0166: 1, H0674: 1, S0364; H0598: 1, H0090: 1, H0040; H0616: 1, H0059: 1, H0100; H0625: 1, H0561: 1, S01432; S0477: 1, H0646: 1, S01432;	1, 5037.2. 1, 1003-0. 1, 5034.2. 1, 50344: 1, H0529: 1, L0763: 1, L0763: 1, L0767: 1, L0768: 1, L0386: 1, L0381: 1, L0803: 1, L0650: 1, L0651: 1, L0529: 1, L0368: 1, L0529: 1, L0368: 1, L4775: 1, L0529: 1, L0368: 1, L4775: 1, L0529: 1, L0368: 1, L4775: 1, L0368: 1,
7, 6, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 7,	2, CO CO CO CO CO CO CO CO CO CO CO CO CO	01	1, SG 1, LOG 1, LOG 1, LOG 1, LOG 1, LOG
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		138079, 138079, 165240, 165240, 165240, 180104, 203740, 219800, 261670, 601649			
		7p13	8		·
1, L0663: 1, L0664: 1, L0665: 1, H0691: 1, L0438: 1, H0519: 1, S0126: 1, H0672: 1, S0330: 1, H0539: 1, S0126: 1, H0552: 1, S0406: 1, H0552: 1, S014: 1, L0742: 1, L0750: 1, L0756: 1, L0777: 1, L0592: 1, L0604: 1, S0242: 1, S0196: 1 and H0352: 1.		AR089: 3, AR316: 2, AR060: 2 L0766: 5, L0803: 5, H0551: 2, L0794: 2, H0435: 2, L0756: 5, L0794: 1, R0282: 1, S0212: 1, S0282: 1, L0534: 1, H0354: 1, H0486: 1, H0014: 1, H0354: 1, R0150: 1, H0647: 1, L0369: 1, L0640: 1, L0771: 1, L0804: 1, L0784: 1, L0792: 1, H0658: 1, L0790: 1, L0799: 1, L0790: 1, L0799: 1, L0799: 1, L0793: 1, And H0543: 1, And H0543: 1	AR060: 2, AR316: 1 L0749: 3, S0354: 2, H0050: and L0602: 1	L0777: 3, L0666: 2, L0750: 2, S0110: 1, S0358: 1, H0251: 1, T0006: 1, L0364: 1, L0659: 1, L0809: 1, H0659: 1, S0378: 1, L0779: 1, L0731: 1 and S0026: 1	AR060: 33, AR316: 31, AR089: 30 L0759: 5, L0783: 4, L0750: 4, L0769: 3, L0774: 3, L0748: 3, L0747: 3, L0731: 3, S0358:
		Pro-26 to Met-35.  A 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	A		Lys-25 to Ser-30. A A A A A A A A A A A A A A A A A A A
2/12	T	5176	5177	5178	5179
191 67	42 - 161	1091 - 1306	183 - 215	531 - 650	594 - 689
2578	8/07	2579	2580	2581	2582
24.1.5	624313	834770	837480	793698	800626
HWI BO67	HWLBUO/	HWLGP26	HWLH031	HWLIL31	HWLJN08
2568	0007	7269	2570	2571	2572

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2, S0408: 2, H0031: 2, L0639: 2, L0775: 2, H0547: 2, L0754: 2, L0775: 2, H0547: 2, L0754: 1, H0341: 1, H0484: 1, S0444: 1, S0360: 1, S0476: 1, H0549: 1, H0550: 1, H0586: 1, H0549: 1, L0622: 1, H0588: 1, H0412: 1, H0100: 1, T0042: 1, H0494: 1, S0144: 1, L0753: 1, L0565: 1, L0565: 1, L0565: 1, L0565: 1, L0576: 1, L057: 1, L057: 1, H0343: 1 and H0707: 1.	AR089: 12, AR316: 11,  AR060: 9  L0758: 9, S0412: 8, L0662: 6, L0749: 6, S0360: 5, H0038: 5, L0751: 5, S0356: 3, H0618: 3, S0051: 3, H0616: 3, L0666: 3, L0743: 3, L0439: 3, L0731: 3, H0295: 2, H0587: 2, H0135: 2, H0591: 2, L0769: 2, L0646: 2, L0774: 2, L0775: 2, L0659: 2, L0774: 2, L0775: 2, L0659: 2, L0665: 2, H0672: 2, S0330: 2, H0539: 2, H0436: 2, L0754: 2, L0588: 2, L0605: 2, H0352: 2, H0539: 1, H0339: 1, H0574: 1, H0241: 1, H0441: 1, H0441: 1, H0286: 1, H033: 1, H0574: 1, S0280: 1, L0021: 1, H0575: 1, H0253: 1, S0010: 1, S0346: 1, H0024: 1, H0031: 1, H033: 1, H0539: 1, H0031: 1, H0588: 1, H0031: 1, H0588: 1, H0031: 1, H0588: 1, H0031: 1, H0588: 1, H0031: 1, H0598: 1, H0598: 1, L0677: 1, L0772:
2, S0408 2, L0777 2, L0748 1, H034 1, S0366 1, H053 1, H010 1, S0144 1, S0144 1, S0428 1, L0657 1, L0657 1, L0657	AR089:
	- 2588 5180
	2526
	RE03 831198
	2573 HWLRE03

1, L0648: 1, L0521: 1, L0767: 1, L0766: 1, L0375: 1, L0378: 1, L0653: 1, L0657: 1, L0783: 1, L0544: 1, L0647: 1, H0435: 1, H0659: 1, H0658: 1, H0670: 1, H0660: 1, H0521: 1, H0696: 1, H0555: 1, H0478: 1, S0037: 1, S0027: 1, L0756: 1, L0777: 1, L0753: 1, H0445: 1, L0596: 1, L0591: 1, S0026: 1 and S0446: 1.	L0771: 5, L0766: 3, H0031: 2, L0794: 2, L0599: 2, H0423: 2, H0265: 1, S0360: 1, T0008: 1, H0042: 1, H0318: 1, H0545: 1, H0014: 1, L0662: 1, L0804: 1, L0805: 1, S0216: 1, L0438: 1, H0659: 1, L0757: 1 and H0352: 1.	AR089: 14, AR316: 11, AR060: 8 L0747: 8, L0758: 8, L0748: 7, L0766: 5, L0776: 4, H0650: 3, H0657: 3, S0420: 3, H0581: 3, H0014: 3, H0670: 3, L0731: 3, S0192: 3, S0340: 2, H0341: 2, H0486: 2, L0770: 2, L0662: 2, L0794: 2, L0770: 2, L0662: 2, L0794: 2, L0743: 2, L0743: 2, H0435: 2, L0770: 1, H053: 2, L0599: 2, L0604: 2, L0361: 2, H0543: 2, H0170: 1, H0583: 1, S0212: 1, H0664: 1, S0358: 1, R0580: 1, S0007: 1, H0587: 1, H0645: 1, L077: 1, H0437: 1, H0642: 1, H0013: 1, H0587: 1, H0642: 1, H0013: 1, H0510: 1, H0251: 1, L0040: 1, H0046: 1, H0123: 1, S0051: 1, H0510:
1, L0648 1, L0766 1, L0653 1, L0544 1, H0655 1, H0667 1, S0027 1, S0027 1, L0753 1, L0753 1, L0753	Pro-27 to Lys-32.  2, L0794 2, H0266 1, H0044 1, H0014 1, L0805 1, H0659	Gly-1 to Lys-6. AR060:  L0747: 7, L0766: 3, H065 3, H0612 4, S0192 5, H043 5, H054 6, H054 1, H064 1, H064 1, H064 1, H064 1, H012
	5181	5182
	377 - 598	1523 - 1564
	602274 2584	793424 25885
	HWTAM38 60	HWTAW58 79
	2574	2575

1, H0375: 1, H0266: 1, H0615: 1, H0553: 1, H0644: 1, H0124: 1, S0036: 1, H0690: 1, H0040: 1, H0551: 1, H0334: 1, H0509: 1, H0633: 1, H0649: 1, S0422: 1, H0659: 1, L0369: 1, L0761: 1, L0768: 1, L0764: 1, L0764: 1, L0768: 1, L0659: 1, L0774: 1, L0657: 1, L0659: 1, L0791: 1, L0530: 1, L0665: 1, H0144: 1, H0702: 1, H0547: 1, S0126: 1, H0555: 1, H065: 1, S0196: 1, L0756: 1, H065: 1, S0194: 1, S0452: 1 and H0352: 1.	H0352: 1	AR089: 14, AR316: 11, AR060: 8 L0439: 14, L0438: 8, S0007: 7, L0803: 7, L0779: 7, L0794: 6, L0766: 6, L0748: 6, L0756: 6, L0731: 6, L0809: 5, L0747: 5, L0749: 5, L0885: 4, L0717: 3, L0805: 3, L0776: 3, L0789: 3, L0751: 3, L0777: 3, L0752: 3, L0759: 3, H0638: 2, S0346: 2, H0024: 2, S0003: 2, L0655: 2, L0770: 2, L0764: 2, L0655: 2, L0770: 2, L0764: 2, L0655: 1, H0170: 1, H0662: 1, S0360: 1, H0193: 1, L0394: 1, S0222: 1, H0486: 1, H0642: 1, H0575: 1, T0110: 1, H0546: 1, H0575: 1, H0615: 1, H0622: 1, H0124: 1, H0163: 1, H0100: 1, H0561: 1, H0132: 1, L0446: 1, L0774: 1, L0662: 1, L0804: 1, L0774:
		Arg-32 to Trp-42.
	5183	5184
	33 - 65	228 - 362
	2586	2587
	603086	855978
	HWTBB42	HWTBC75
	2576	2577

1, L0775: 1, L0657: 1, L0636: 1, L0367: 1, L0643: 1, H0520: 1, H0547: 1, S0126: 1, H0690: 1, H0658: 1, H0648: 1, S0328: 1, L0750: 1, L0780: 1, L0757: 1, H0445: 1, L0757: 1, H0645: 1, L0599: 1, L0593: 1, L0595: 1, L0566: 1, H0668: 1, H0665: 1, S0242: 1 and H0352: 1.	AR060: 22, AR316: 19, AR089: 15 H0352: 2 and L0352: 1.	AR089: 15, AR316: 13, AR060: 11 H0556: 8, L0439: 7, L0779: 7, L0766: 6, L0803: 6, H0013: 5, L0758: 4, H0624: 3, H0575: 3, L0774: 3, L0748: 3, H0575: 2, L0761: 2, L0804: 2, L0805: 2, L0761: 2, L0804: 2, L0805: 2, L0751: 2, L0747: 2, L0509: 2, L0751: 2, L0777: 2, L0509: 2, L0756: 2, L0777: 2, L0509: 2, L0756: 2, L0777: 2, L0509: 1, L0615: 1, H0680: 1, H0717: 1, S0134: 1, H0650: 1, H0577: 1, H0656: 1, S0418: 1, S0420: 1, H0656: 1, R0194: 1, H0080: 1, H068: 1, H0194: 1, H0080: 1, H0178: 1, H0563: 1, H0644: 1, H0178: 1, H0563: 1, H0640: 1, H0178: 1, H050: 1, H0591: 1, H0178: 1, H050: 1, H0591: 1, H0178: 1, H050: 1, H0591: 1, H0024: 1, H0194: 1, H0640: 1, H0077: 1, H0641: 1, L0761: 1, H0087: 1, H0690: 1, L0761: 1, L0769: 1, L0800: 1, L0764:
	Pro-27 to Tyr-33.	A A L L L L L L L L L L L L L L L L L L
	5185	5186
	197 - 298	2270 - 2386
	2588	2589
	637736	791720
	HWTB125	HWTBL86
,	2578	2579

1, L0521: 1, L0794: 1, L0649: 1, L0381: 1, L0775: 1, L0651: 1, L0378: 1, L0776: 1, L0655: 1, L0661: 1, L0657: 1, L0659: 1, L0789: 1, L0790: 1, L4501: 1, L0789: 1, L0790: 1, L4501: 1, L0666: 1, S0126: 1, H0682: 1, H0658: 1, H0670: 1, S0328: 1, S0330: 1, H0521: 1, H0696: 1, S0404: 1, H0436: 1, S3012: 1, S0404: 1, L0743: 1, L0740: 1, L0754: 1, L0752: 1, L0759: 1, H0667: 1, S0194: 1, H0543: 1, H06423: 1 and S0424: 1.	AR204: 220, AR205: 174, AR060: 161, AR244: 155, AR104: 155, AR243: 149, AR232: 149, AR192: 147, AR186: 142, AR185: 137, AR312: 134, AR274: 133, AR206: 132, AR246: 130, AR206: 132, AR246: 130, AR206: 132, AR299: 127, AR039: 125, AR177: 125, AR194: 120, AR313: 119, AR207: 113, AR096: 112, AR291: 114, AR295: 114, AR291: 114, AR295: 110, AR285: 106, AR286: 100, AR198: 108, AR286: 100, AR209: 101, AR266: 100, AR209: 101, AR266: 100, AR207: 99, AR184: 97, AR271: 97, AR233: 96, AR247: 94, AR256: 94, AR247: 94, AR298: 89, AR247: 94, AR298: 89,
	Arg-48 to Val-53, AF Ala-57 to Ala-69. AF Ala-57 to Ala-69. AF AF AF AF AF AF AF AF AF AF AF AF AF
	261 - 1076 5187
	732187 2590
	2580 HWTBX66

AR275: 86, AR259: 86, AR055: 85, AR053: 83, AR238: 82, AR240: 82, AR310: 80, AR218: 79, AR293: 75, AR229: 72, AR290: 70, AR234: 70, AR290: 70, AR234: 70, AR249: 68, AR267: 69, AR249: 68, AR267: 64, AR270: 65, AR277: 64, AR270: 60, AR269: 58, AR280: 56, AR315: 54, AR281: 50, AR314: 49, AR281: 50, AR314: 49, AR231: 49 S0468: 1, L0770: 1 and H0352: 1.	H0583: 1, H0030: 1, H0135: , L0773: 1 and S0136: 1.	H0583: 1 AR060: 3, AR316: 2,	AR089: 2 H0583: 1	AR060: 5, AR316: 4, AR089: 4 S0282: 10, H0594: 6, H0596: 5, H0592: 3, S0260: 3, H0595: 3, H0600: 2, H0041: 2, H0105: 2, S0388: 2, H0316: 2, H0598: 2, S0344: 2, S0390: 2, S0218: 1, H0411: 1, S0280: 1, H0590: 1, H0559: 1, S0194: 1 and S0042: 1.	H0586: 1, H0036: 1, H0041: 1, H0271: 1, H0622: 1, H0488: 1, H0509: 1 and S0434: 1.
<u> </u>		A	<b>A</b>	4 4	Glu-32 to Arg-38, Val-46 to Val-61.
	5188	5189 5190		5191	5192
	197 - 313	341 - 397 461 - 556	,	166 - 279	261 - 467
	2591	2592 2593		2594	2595
	762843	897736 696654		853561	898367
		HYAAD61 HYACC21		HYBAP75	HYBAQ24
	2581	2582 2583		2584	2585

AR089: 144, AR316: 101,	AR060: 61	H0041: 1	H0041: 1
Arg-21 to Lys-27.			
5193			5194
116 - 268			151 - 219
2596			2597
658736			533784
2586   HYBAW56   658736			2587 HYBBD81
2586			2587

[78] The first column in Table 1B provides the gene number in the application corresponding to the clone identifier. The second column in Table 1B provides a unique "Clone ID NO:Z" for a cDNA clone related to each contig sequence disclosed in Table 1B. This clone ID references the cDNA clone which contains at least the 5' most sequence of the assembled contig and at least a portion of SEQ ID NO:X was determined by directly sequencing the referenced clone. The reference clone may have more sequence than described in the sequence listing or the clone may have less. In the vast majority of cases, however, the clone is believed to encode a full-length polypeptide. In the case where a clone is not full-length, a full-length cDNA can be obtained by methods described elsewhere herein.

[79] The third column in Table 1B provides a unique "Contig ID" identification for each contig sequence. The fourth column provides the "SEQ ID NO:" identifier for each of the contig polynucleotide sequences disclosed in Table 1B. The fifth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence "SEQ ID NO:X" that delineate the preferred open reading frame (ORF) shown in the sequence listing and referenced in Table 1B, column 6, as SEQ ID NO:Y. Where the nucleotide position number "To" is lower than the nucleotide position number "From", the preferred ORF is the reverse complement of the referenced polynucleotide sequence.

[80] The sixth column in Table 1B provides the corresponding SEQ ID NO:Y for the polypeptide sequence encoded by the preferred ORF delineated in column 5. In one embodiment, the invention provides an amino acid sequence comprising, or alternatively consisting of, a polypeptide encoded by the portion of SEQ ID NO:X delineated by "ORF (From-To)". Also provided are polynucleotides encoding such amino acid sequences and the complementary strand thereto.

[81] Column 7 in Table 1B lists residues comprising epitopes contained in the polypeptides encoded by the preferred ORF (SEQ ID NO:Y), as predicted using the algorithm of Jameson and Wolf, (1988) Comp. Appl. Biosci. 4:181-186. The Jameson-

Wolf antigenic analysis was performed using the computer program PROTEAN (Version 3.11 for the Power MacIntosh, DNASTAR, Inc., 1228 South Park Street Madison, WI). In specific embodiments, polypeptides of the invention comprise, or alternatively consist of, at least one, two, three, four, five or more of the predicted epitopes as described in Table 1B. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly.

[82] Column 8, in Table 1B, provides an expression profile and library code: count for each of the contig sequences (SEQ ID NO:X) disclosed in Table 1B, which can routinely be combined with the information provided in Table 4 and used to determine the tissues, cells, and/or cell line libraries which predominantly express the polynucleotides of the invention. The first number in column 8 (preceding the colon), represents the tissue/cell source identifier code corresponding to the code and description provided in Table 4. For those identifier codes in which the first two letters are not "AR", the second number in column 8 (following the colon) represents the number of times a sequence corresponding to the reference polynucleotide sequence was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array technology. Utilizing this technology, cDNAs were amplified by PCR and then transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array. cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of <sup>33</sup>P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding

polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression.

[83] Column 9 in Table 1B provides a chromosomal map location for certain polynucleotides of the invention. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Each sequence in the UniGene database is assigned to a "cluster"; all of the ESTs, cDNAs, and STSs in a cluster are believed to be derived from a single gene. Chromosomal mapping data is often available for one or more sequence(s) in a UniGene cluster; this data (if consistent) is then applied to the cluster as a whole. Thus, it is possible to infer the chromosomal location of a new polynucleotide sequence by determining its identity with a mapped UniGene cluster.

[84] A modified version of the computer program BLASTN (Altshul, et al., J. Mol. Biol. 215:403-410 (1990), and Gish, and States, Nat. Genet. 3:266-272) (1993) was used to search the UniGene database for EST or cDNA sequences that contain exact or near-exact matches to a polynucleotide sequence of the invention (the 'Query'). A sequence from the UniGene database (the 'Subject') was said to be an exact match if it contained a segment of 50 nucleotides in length such that 48 of those nucleotides were in the same order as found in the Query sequence. If all of the matches that met this criteria were in the same UniGene cluster, and mapping data was available for this cluster, it is indicated in Table 1B under the heading "Cytologic Band". Where a cluster had been further localized to a distinct cytologic band, that band is disclosed; where no banding information was available, but the gene had been localized to a single chromosome, the chromosome is disclosed.

[85] Once a presumptive chromosomal location was determined for a polynucleotide of the invention, an associated disease locus was identified by comparison with a database of diseases which have been experimentally associated with genetic loci. The database used was the Morbid Map, derived from OMIM<sup>TM</sup> ("Online Mendelian Inheritance in Man"; McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National Library of Medicine (Bethesda, MD) 2000; World Wide Web URL: http://www.ncbi.nlm.nih.gov/omim/). If the putative chromosomal location of a polynucleotide of the invention (Query sequence)

was associated with a disease in the Morbid Map database, an OMIM reference identification number was noted in column 10, Table 1B, labelled "OMIM Disease Reference(s). Table 5 is a key to the OMIM reference identification numbers (column 1), and provides a description of the associated disease in Column 2.

[86] Table 1C summarizes additional polynucleotides encompassed by the invention (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) contig nucleotide sequence identifiers (SEQ ID NO:X)), and genomic sequences (SEQ ID NO:B). The first column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence. The second column provides the sequence identifier, "SEQ ID NO:X", for each contig sequence. The third column provides a unique contig identifier, "Contig ID:" for each contig sequence. The fourth column, provides a BAC identifier "BAC ID NO:A" for the BAC clone referenced in the corresponding row of the table. The fifth column provides the nucleotide sequence identifier, "SEQ ID NO:B" for a fragment of the BAC clone identified in column four of the corresponding row of the table. The sixth column, "Exon From-To", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:B which delineate certain polynucleotides of the invention that are also exemplary members of polynucleotide sequences that encode polypeptides of the invention (e.g., polypeptides containing amino acid sequences encoded by the polynucleotide sequences delineated in column six, and fragments and variants thereof).

## TABLE 1C

Clone ID No:Z	SEQ ID NO:X	CONTIG ID:	BAC ID: A	SEQ ID NO:B	EXON From-To
Н2СВН91	12	826669	AF260012	5205	1-624
IIZ CDII) I	1.2	02000)	AI 200012	3203	4141-4362
					4976-5265
					5433-5573
					5783-5890
					6096-6635
					7710-8129
H2CBH91	12	826669	AF260012	5206	8177-10172
112CD1191	12	020009	AF200012	3200	1-449
H2CBH91	12	826669	AF260012	5207	512-986
112CD1191	12	820009	AF200012	3207	1-80
					527-631
					1970-2099
					2509-2582
				1	2740-2802
					3491-3728
					3976-4184
					4306-4978
				.	5841-6002
HOLDBOO	14	650665	1.0015050	1	7317-8173
H2LBB09	14	658667	AC015863	5208	1-103
					2561-2824
					4532-4708
					5840-7609
H2LBB09	15	830636	AC015863	5209	1-103
					2561-2824
					4532-4708
					5840-7609
H2MAC63	16	610045	AC019122	5210	1-63
			·		390-1660
					3921-3948
H2MAC63	16	610045	AC019122	5211	1-331
H2MAC63	16	610045	AC019122	5212	1-450
					539-708
					713-835
H2MBA76	17	826161	AP001822	5213	1-69
					608-1096
					1251-2446
H2MBA76	17	826161	AC020997	5214	1-201
					1064-1126
					1665-2153
					2308-3502
H2MBA76	17	826161	AP001822	5215	1-71
	_,		111001022		929-1051
					1659-1798
H2MBA76	17	826161	AC020997	5216	1-1329
H2MBA76	17	826161	AC020997	5217	1-1329
	**	020101	110020991	3211	
					930-1052
H2MBF60	18	695714	AC015470	5310	1660-1799
112IVIDFOU	10	093/14	AC015470	5218	1-470
HOMBECO	10	(05714	4.0015.450	15010	850-1865
H2MBF60	18	695714	AC015470	5219	1-244
H6BSM88	19	609832	AL355520	5220	1-39
					3561-3618
					7841-7949

H6BSM88	1275 11270
H6BSM88   19	0375-11270
H6BSM88   19	1589-11678
H6BSM88   19	5048-15690
H6BSM88	5028-16134
H6BSM88   19	3215-18476
H6BSM88   19	0699-20733
H6BSM88	3616-23861
H6BSM88	0319-35764
H6BSM88	1-386
H6EEA48       20       847111       AC011479       5223         H6EEA48       20       847111       AC011479       5224         H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	1-437
H6EEA48 20 847111 AC011479 5224 H6EEU40 23 757048 AC026285 5225 H6EEU40 23 757048 AC023920 5226 H6EEU40 23 757048 AC026285 5227 H6EEU40 23 757048 AC023920 5228 H7TMB95 25 597427 AL133381 5229 H7TMB95 25 597427 AL133385 5230 HACAD42 27 634372 AC018512 5231 HACAD42 27 634372 AC02305 5232 HACAD42 27 634372 AC02518 5233 HACAD42 27 634372 AC02518 5233 HACAD42 27 634372 AC02518 5233 HACAD42 27 634372 AC016135 5234 HACAD42 27 634372 AC016135 5234 HACAD40 33 745367 AC011498 5235	1-252
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	298-434
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	597-678
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	805-867
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	1903-2229
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	2668-2872
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	3606-3709
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	4150-4555
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H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	4973-5096
H6EEU40       23       757048       AC026285       5225         H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC02305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	5167-6102
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H6EEU40       23       757048       AC023920       5226         H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC022305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	1-246
H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC022305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	1-414
H6EEU40       23       757048       AC026285       5227         H6EEU40       23       757048       AC023920       5228         H7TMB95       25       597427       AL133381       5229         H7TMB95       25       597427       AL133385       5230         HACAD42       27       634372       AC018512       5231         HACAD42       27       634372       AC022305       5232         HACAD42       27       634372       AC002518       5233         HACAD42       27       634372       AC016135       5234         HADAB60       33       745367       AC011498       5235	440-1918
H6EEU40 23 757048 AC023920 5228  H7TMB95 25 597427 AL133381 5229  H7TMB95 25 597427 AL133385 5230  HACAD42 27 634372 AC018512 5231  HACAD42 27 634372 AC02305 5232  HACAD42 27 634372 AC002518 5233  HACAD42 27 634372 AC016135 5234  HACAD42 37 634372 AC016135 5234  HACAD460 33 745367 AC011498 5235	1-416
H6EEU40 23 757048 AC023920 5228  H7TMB95 25 597427 AL133381 5229  H7TMB95 25 597427 AL133385 5230  HACAD42 27 634372 AC018512 5231  HACAD42 27 634372 AC02305 5232  HACAD42 27 634372 AC002518 5233  HACAD42 27 634372 AC016135 5234  HACAD42 37 634372 AC016135 5234  HACAD42 37 AC011498 5235	442-1920
H7TMB95 25 597427 AL133381 5229 H7TMB95 25 597427 AL133385 5230 HACAD42 27 634372 AC018512 5231 HACAD42 27 634372 AC022305 5232 HACAD42 27 634372 AC002518 5233 HACAD42 27 634372 AC016135 5234 HADAB60 33 745367 AC011498 5235	1-1458
H7TMB95 25 597427 AL133381 5229 H7TMB95 25 597427 AL133385 5230 HACAD42 27 634372 AC018512 5231 HACAD42 27 634372 AC022305 5232 HACAD42 27 634372 AC002518 5233 HACAD42 27 634372 AC016135 5234 HADAB60 33 745367 AC011498 5235	1729-1836
H7TMB95 25 597427 AL133385 5230 HACAD42 27 634372 AC018512 5231 HACAD42 27 634372 AC022305 5232 HACAD42 27 634372 AC002518 5233 HACAD42 27 634372 AC016135 5234 HADAB60 33 745367 AC011498 5235	1-1459
H7TMB95 25 597427 AL133385 5230 HACAD42 27 634372 AC018512 5231 HACAD42 27 634372 AC022305 5232 HACAD42 27 634372 AC002518 5233 HACAD42 27 634372 AC016135 5234 HADAB60 33 745367 AC011498 5235	1730-1837
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	1-3168
	3709-3896
	4647-4876
	5003-5294
	5443-5552
	5836-6190
	6334-6407
	7922-8433
	9129-9255
	9889-10202
1 1 1 1 1 1 1	0311-10385
	0503-10664
	2531-12600
1 1 1 1	2682-12801
1 1 1 1	5489-15696
HADAB60 33 745367 AC011498 5236	1-165
	3986-4070
	4146-4408
HADAB60 33 745367 AC011498 5237	1-385
HADAM31 34 604500 AC001084 5238	1-833
HADAM31 34 604500 AC008670 5239	1-833
HADAM31 34 604500 AC008670 5240	1-358
HADCL19 35 599065 AC027264 5241	1-147
HADCL19 35 599065 AL353663 5242	1-141

HADCL19 HADCL19	35	599065	AC023583	5243	1-183
	35	599065	AC027584	5244	1-162
HADCLIS	35	599065	AC023309	5245	1-193
	35	599065	AC068682	5246	1-153
HADCL19		599065	AL356423	5247	1-203
HADCL19	35	599065	AC016142	5248	1-150
HADCL19	35		AC010142 AC023864	5249	1-130
HADCL19	35	599065		5250	1-142
HADCL19	35	599065	AC021466		1-103
HADCL19	35	599065	AC000360	5251	1-368
HADCL19	35	599065	AC027584	5252	1-308
HADCL19	35	599065	AC023864	5253	1590-4704
HADDC04	37	601695	AC025086	5254	1-2523
HADDC04	37	601695	AC036213	5255	1-2523
HADDC04	37	601695	AC025086	5256	1-186
HADDP23	38	847115	AL353606	5257	1-931
HADDP23	38	847115	AC022252	5258	1-931
HADDP23	38	847115	AL365360	5259	1-931
HADDR24	40	664477	AF047825	5260	1-1062
HADDR24	40	664477	AC005103	5261	1-1062
HADDR24	40	664477	AF047825	5262	1-124
HADDR24	40	664477	AF047825	5263	1-512
HADDR24	40	664477	AC005103	5264	1-512
HADDR24		664477	AC005103	5265	1-128
	40	607615	AL139187	5266	1-267
HADET62	41	00/013	AL139107	3200	881-1219
					3142-3948
					5639-5823
TIADETCO	111	607615	AL139187	5267	1-355
HADET62	41	704584	AC006159	5268	1-813
HADEY22	43	704364	AC000139	3208	1374-1711
					3195-3417
					6590-6718
	1				7485-7921
		ļ		ļ	9403-9953
					11546-12050
	1				14701-15135
					19748-20021
				ļ	20944-21554
					25746-26009
					26183-26508
					33951-36192
HADEY22	44	861628	AC006159	5269	1-813
111111111111111111111111111111111111111	' '	"""			1374-1711
					3195-3417
					6590-6718
					7485-7921
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,					9403-9953
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,					11546-12050
,					11546-12050 14701-15135
					11546-12050 14701-15135 19748-20021
					11546-12050 14701-15135 19748-20021 20944-21554 25746-26009
					11546-12050 14701-15135 19748-20021 20944-21554 25746-26009 26183-26508
HADER84	45	668229	AC034117	5270	11546-12050 14701-15135 19748-20021 20944-21554 25746-26009 26183-26508 33951-36192
HADFB84	45	668229 668229	AC034117 AC034117	5270 5271	11546-12050 14701-15135 19748-20021 20944-21554 25746-26009 26183-26508 33951-36192
HADFB84	45	668229	AC034117	5271	11546-12050 14701-15135 19748-20021 20944-21554 25746-26009 26183-26508 33951-36192 1-3119 1-216
					11546-12050 14701-15135 19748-20021 20944-21554 25746-26009 26183-26508 33951-36192

HADFK11	48	637485	Z97353	5275	1-286
HADFT44	49	793100	AC010789	5276	1-67
					205-268
					557-1982
HADFT44	49	793100	AC016774	5277	1-67
					205-268
					557-1982
HADFT44	49	793100	AC010789	5278	1-302
HADFT44	49	793100	AC010789	5279	1-544
HADFT44	49	793100	AC016774	5280	1-544
HADFT44	49	793100	AC016774	5281	1-302
HADFW20	50	599066	AC009314	5282	1-227
111101 1120		333000	110007011		1999-2922
					2929-3376
					3543-4652
					5475-5580
		•			5690-5858
					8208-8577
				ļ	10545-10753
				1	11835-12027
					13238-14161
HADFW20	50	599066	AC009314	5283	1-767
HADFW20	50	599066	AC009314	5284	1-473
	51	741054	AF139813	5285	1-1204
HADFX10		741054	AC004228	5286	1-1204
HADFX10	51		AC004228 AC003025	5287	1-1204
HADFX10	51	741054		5288	1-1204
HADFX10	51	741054	AF139813		
HADFX10	51	741054	AC004228	5289	1-332
HADFX10	51	741054	AC003025	5290	1-332
HADFY80	52	654831	AC016498	5291	1-33
					3823-3948
					4874-5669
HADFY80	52	654831	AC026740	5292	1-33
					3772-3897
					4823-5339
HADFY80	52	654831	AC016498	5293	1-451
HADFY80	52	654831	AC026740	5294	1-115
					765-1186
HADMA77	54	783049	AC007944	5295	1-3350
HADMA77	54	783049	AC018656	5296	1-3349
HADMA77	54	783049	AC021874	5297	1-3351
					4529-4959
					6110-6438
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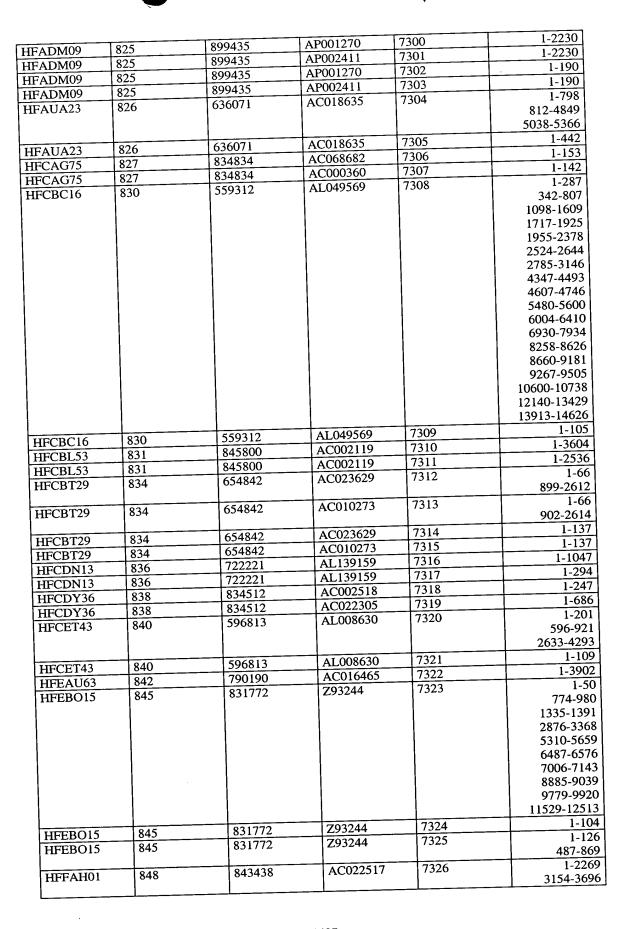
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